

10/593545  
IAPG/ROC/PCT/PTO 20 SEP 2005

PCT-US2005-009928\_Sequence Listing.txt as filed 10-3-05 in PCT  
SEQUENCE LISTING

<110> Rockefeller University

<120> LYTIC ENZYMES AND SPORE SURFACE ANTIGEN FOR DETECTION AND TREATMENT OF  
BACILLUS ANTHRACIS BACTERIA AND SPORES

<130> 12157/17

<140> PCT/US 2005/009928

<141> 2005-03-23

<150> 60/555,916

<151> 2004-03-24

<160> 109

<170> PatentIn version 3.1

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<212> DNA

<213> Bacillus anthracis

<400> 1

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<213> *Bacillus anthracis*

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Leu Thr Ala Ala Gln Lys Lys Glu Phe Asp Thr Leu Ala Ala Glu Leu  
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Val Arg Leu Lys Ile Phe Ser Asn Leu Asp Val Asp Ser Leu Ala Arg  
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Tyr Val Asp Ser Lys Asp Gln Tyr Ile Lys Met Val Arg Leu Leu Arg  
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Lys Thr Lys Pro Ser Asp Asp Phe Lys Leu Tyr Ser Gln Met Gln Arg  
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Ser Lys Asn Leu Leu Phe Asn Glu Cys Arg Ser Ser Ala Ser Asp Leu  
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Gly Leu Thr Ile Thr Ser Arg Leu Lys Leu Val Ile Pro Glu Val Asp  
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Leu Thr Ala Ala Gln Lys Lys Glu Phe Asp Thr Leu Ala Ala Glu Leu  
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Val Arg Leu Lys Ile Phe Ser Asn Leu Asp Val Asp Ser Leu Ala Arg  
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Tyr Val Asp Ser Lys Asp Gln Tyr Ile Lys Met Val Arg Leu Leu Arg  
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Lys Thr Lys Pro Ser Asp Asp Phe Lys Leu Tyr Ser Gln Met Gln Arg  
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Ser Lys Asn Leu Leu Phe Asn Glu Cys Arg Ser Ser Ala Ser Asp Leu  
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Gly Leu Thr Ile Thr Ser Arg Leu Lys Leu Val Ile Pro Glu Val Asp  
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&lt;213&gt; Bacillus anthracis

&lt;400&gt; 5

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Phe Ile Arg Asp Tyr Glu Glu Cys Gln Ser Glu Asp Ser Pro Phe Tyr  
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Phe Asp Gly Glu Ile Ala Glu Asp Phe Tyr Trp Phe Ala Lys Glu Phe  
 50 55 60

Lys His Val Glu Gly Ile Leu Ala Gly Glu Ser Val Glu Leu Thr Asp  
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Phe Gln Leu Phe Leu Ala Ala Asn Ile Phe Gly Phe Lys Lys Lys Ile  
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Asn Gly Ala Arg Arg Phe Arg Lys Val Phe Ile Gln Leu Ala Arg Lys  
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Asn Ala Lys Ser Gln Phe Leu Ala Ile Val Ala Ala Phe Cys Thr Phe  
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Leu Gly Asp Glu Lys Gln Arg Ala Tyr Ile Ala Gly Trp Thr Arg Asp  
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Gln Ser Ser Glu Val Tyr Glu Ala Val Lys Thr Gly Ile Ser Ser Ser  
 145 150 155 160

Glu Leu Leu Glu Gly Lys Trp Lys Glu Ala Tyr Ser Thr Ile Glu Ile  
 165 170 175

Phe Lys Asn Gly Ser Val Val Val Pro Leu Ser Lys Glu Ala Arg Lys  
 180 185 190

Thr Gly Asp Gly Lys Asn Pro Ser Leu Gly Ile Val Asp Glu Tyr His  
 195 200 205

Ala His Glu Thr Asp Glu Ile Tyr Asp Val Leu Ser Ser Gly Met Val  
 210 215 220

Ala Arg Lys Glu Pro Leu Met Phe Ile Ile Thr Thr Ala Gly Phe Asp  
 225 230 235 240

Leu Ser Arg Pro Cys Tyr Arg Glu Tyr Glu Tyr Val Ser Asp Ile Leu  
 245 250 255

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Asp Pro Ser Lys Asn Val Glu Asn Asp Asp Tyr Phe Val Met Ile Cys  
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Glu Leu Glu Lys Asn Asp Asp Ile Lys Asp Glu Ser Asn Trp Ile Lys  
275 280 285

Ala Asn Pro Ile Val Ala Thr Tyr Glu Glu Gly Leu Glu Gly Ile Arg  
290 295 300

Ser Asp Leu Lys Val Ala Leu Asp Arg Pro Glu Lys Met Arg Ala Phe  
305 310 315 320

Leu Thr Lys Asn Met Asn Ile Trp Val Asp Lys Lys Asp Asn Gly Tyr  
325 330 335

Met Asp Met Ser Lys Trp Gln Lys Cys Glu Val Asp Thr Phe Asp Phe  
340 345 350

Ser Gly Ala Thr Leu Trp Ile Gly Gly Asp Leu Ser Met Thr Thr Asp  
355 360 365

Leu Thr Ser Val Gly Trp Val Gly Met Asp Asp Glu Gly Asp Phe Ile  
370 375 380

Val Gly Gln His Ser Phe Met Pro Glu Ala Arg Leu Lys Glu Lys Met  
385 390 395 400

Ala Ile Asp Lys Val Arg Tyr Asp Leu Trp Ala Glu Gln Gly Tyr Leu  
405 410 415

Thr Leu Thr Pro Gly Glu Met Val Asp Tyr Thr Ile Val Glu Ser Trp  
420 425 430

Ile Glu Asn Phe Ser Lys Asp Lys Glu Ile Gln Glu Phe Asp Tyr Asp  
435 440 445

Lys Trp Asn Ala Leu His Leu Ala Gln Asn Leu Glu Asn Lys Gly Phe  
450 455 460

Val Cys Val Glu Ile Pro Gln Arg Ile Ala Asn Leu Ser Ile Pro Thr  
465 470 475 480

Lys Asn Phe Arg Glu Lys Val Tyr Glu Lys Lys Val Lys His Asn Gly  
485 490 495

Asp Pro Val Leu Phe Trp Ala Leu Asn Asn Ala Val Val Lys Met Asp  
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Asp Gln Glu Asn Ile Met Ile Ser Lys Lys Ile Ser Lys Asn Arg Ile



Asp Pro Ala Ala Ala Val Leu Asn Ala Phe Ser Arg Ala Met Tyr Gly  
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Gly Lys Leu Trp Asn  
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<212> PRT  
<213> Bacillus anthracis

<400> 6

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Phe Ile Arg Asp Tyr Glu Glu Cys Gln Ser Glu Asp Ser Pro Phe Tyr  
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Phe Asp Gly Glu Ile Ala Glu Asp Phe Tyr Trp Phe Ala Lys Glu Phe  
50 55 60

Lys His Val Glu Gly Ile Leu Ala Gly Glu Ser Val Glu Leu Thr Asp  
65 70 75 80

Phe Gln Leu Phe Leu Ala Ala Asn Ile Phe Gly Phe Lys Lys Lys Ile  
85 90 95

Asn Gly Ala Arg Arg Phe Arg Lys Val Phe Ile Gln Leu Ala Arg Lys  
100 105 110

Asn Ala Lys Ser Gln Phe Leu Ala Ile Val Ala Ala Phe Cys Thr Phe  
115 120 125

Leu Gly Asp Glu Lys Gln Arg Ala Tyr Ile Ala Gly Trp Thr Arg Asp  
130 135 140

Gln Ser Ser Glu Val Tyr Glu Ala Val Lys Thr Gly Ile Ser Ser Ser  
145 150 155 160

Glu Leu Leu Glu Gly Lys Trp Lys Glu Ala Tyr Ser Thr Ile Glu Ile  
165 170 175

Phe Lys Asn Gly Ser Val Val Val Pro Leu Ser Lys Glu Ala Arg Lys  
Page 44

Thr Gly Asp Gly Lys Asn Pro Ser Leu Gly Ile Val Asp Glu Tyr His  
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Ala His Glu Thr Asp Glu Ile Tyr Asp Val Leu Ser Ser Gly Met Val  
210 215 220

Ala Arg Lys Glu Pro Leu Met Phe Ile Ile Thr Thr Ala Gly Phe Asp  
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Leu Ser Arg Pro Cys Tyr Arg Glu Tyr Glu Tyr Val Ser Asp Ile Leu  
245 250 255

Asp Pro Ser Lys Asn Val Glu Asn Asp Asp Tyr Phe Val Met Ile Cys  
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Glu Leu Glu Lys Asn Asp Asp Ile Lys Asp Glu Ser Asn Trp Ile Lys  
275 280 285

Ala Asn Pro Ile Val Ala Thr Tyr Glu Glu Gly Leu Glu Gly Ile Arg  
290 295 300

Ser Asp Leu Lys Val Ala Leu Asp Arg Pro Glu Lys Met Arg Ala Phe  
305 310 315 320

Leu Thr Lys Asn Met Asn Ile Trp Val Asp Lys Lys Asp Asn Gly Tyr  
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Met Asp Met Ser Lys Trp Gln Lys Cys Glu Val Asp Thr Phe Asp Phe  
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Ser Gly Ala Thr Leu Trp Ile Gly Gly Asp Leu Ser Met Thr Thr Asp  
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Leu Thr Ser Val Gly Trp Val Gly Met Asp Asp Glu Gly Asp Phe Ile  
370 375 380

Val Gly Gln His Ser Phe Met Pro Glu Ala Arg Leu Lys Glu Lys Met  
385 390 395 400

Ala Ile Asp Lys Val Arg Tyr Asp Leu Trp Ala Glu Gln Gly Tyr Leu  
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Thr Leu Thr Pro Gly Glu Met Val Asp Tyr Thr Ile Val Glu Ser Trp  
420 425 430

Ile Glu Asn Phe Ser Lys Asp Lys Glu Ile Gln Glu Phe Asp Tyr Asp  
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Lys Trp Asn Ala Leu His Leu Ala Gln Asn Leu Glu Asn Lys Gly Phe  
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Val Cys Val Glu Ile Pro Gln Arg Ile Ala Asn Leu Ser Ile Pro Thr  
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Lys Asn Phe Arg Glu Lys Val Tyr Glu Lys Lys Val Lys His Asn Gly  
485 490 495

Asp Pro Val Leu Phe Trp Ala Leu Asn Asn Ala Val Val Lys Met Asp  
500 505 510

Asp Gln Glu Asn Ile Met Ile Ser Lys Lys Ile Ser Lys Asn Arg Ile  
515 520 525

Asp Pro Ala Ala Ala Val Leu Asn Ala Phe Ser Arg Ala Met Tyr Gly  
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Gly Lys Leu Trp Asn  
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Glu Trp Leu Gly Ile Ser Pro Ser Thr Ile Ser Val Lys Gly Lys Asn  
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Ala Leu Lys Val Ala Thr Val Phe Ala Cys Ile Lys Ile Leu Ser Glu  
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Ser Val Ser Lys Leu Pro Leu Lys Ile Tyr Gln Glu Asp Glu Tyr Gly  
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Ile Gln Arg Gly Thr Lys His Tyr Leu Asn Asn Leu Leu Arg Leu Arg  
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Pro Asn Pro Tyr Met Ser Ser Met Asn Phe Phe Gly Ser Leu Glu Ala  
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 Gln Lys Asn Leu Tyr Gly Asn Ser Tyr Ala Asn Ile Glu Phe Asp Arg  
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Val Tyr Ile Asp Asp Val Gly Leu Leu Asn Ser Lys Thr Lys Met Trp  
 145 150 155 160

Tyr Val Val Asn Thr Gly Gly Gln Gln Arg Val Leu Lys Pro Glu Glu  
 165 170 175

Ile Leu His Phe Lys Asn Gly Ile Thr Leu Asp Gly Leu Val Gly Val  
 180 185 190

Pro Thr Met Glu Tyr Leu Lys Ser Thr Leu Glu Asn Ser Ala Ser Ala  
 195 200 205

Asp Lys Phe Ile Asn Asn Phe Tyr Lys Gln Gly Leu Gln Val Lys Gly  
 210 215 220

Leu Val Gln Tyr Val Gly Asp Leu Asn Glu Asp Ala Lys Lys Val Phe  
 225 230 235 240

Arg Glu Asn Phe Glu Ser Met Ser Ser Gly Leu Gln Asn Ser His Arg  
 245 250 255

Ile Ala Leu Met Pro Val Gly Tyr Gln Phe Gln Pro Ile Ser Leu Asn  
 260 265 270

Met Ser Asp Ala Gln Phe Leu Glu Asn Thr Glu Leu Thr Ile Arg Gln  
 275 280 285

Ile Ala Thr Ala Phe Gly Ile Lys Met His Gln Leu Asn Asp Leu Ser  
 290 295 300

Lys Ala Thr Leu Asn Asn Ile Glu Gln Gln Gln Gln Gln Phe Tyr Thr  
 305 310 315 320

Asp Thr Leu Gln Ala Thr Leu Thr Met Tyr Glu Gln Glu Met Thr Tyr  
 325 330 335

Lys Leu Phe Leu Asp Ser Glu Leu Asp Lys Gly Phe Tyr Ser Lys Phe  
 340 345 350

Asn Val Asp Ala Ile Leu Arg Ala Asp Ile Lys Thr Arg Tyr Glu Ala  
 355 360 365

Tyr Arg Thr Gly Ile Gln Gly Gly Phe Leu Lys Pro Asn Glu Ala Arg  
 370 375 380

PCT-US2005-009928\_Sequence Listing.txt as filed 10-3-05 in PCT

Ser Lys Glu Asp Leu Pro Pro Glu Ala Gly Gly Asp Arg Leu Leu Val  
385 390 395 400

Asn Gly Asn Met Leu Pro Ile Asp Met Ala Gly Gln Ala Tyr Leu Lys  
405 410 415

Gly Gly Asp Thr Asn Gly Glu Val Ser Lys Glu Gly Asn Glu Gly Asn  
420 425 430

<210> 8  
<211> 432  
<212> PRT  
<213> Bacillus anthracis

<400> 8

Val Lys Ile Val Asp Ser Val Lys Lys Phe Phe Asn Phe Glu Lys Arg  
1 5 10 15

Gln Thr Ser Gln Val Ile Glu Leu Asn Lys Asp Asp Glu Lys Leu Leu  
20 25 30

Glu Trp Leu Gly Ile Ser Pro Ser Thr Ile Ser Val Lys Gly Lys Asn  
35 40 45

Ala Leu Lys Val Ala Thr Val Phe Ala Cys Ile Lys Ile Leu Ser Glu  
50 55 60

Ser Val Ser Lys Leu Pro Leu Lys Ile Tyr Gln Glu Asp Glu Tyr Gly  
65 70 75 80

Ile Gln Arg Gly Thr Lys His Tyr Leu Asn Asn Leu Leu Arg Leu Arg  
85 90 95

Pro Asn Pro Tyr Met Ser Ser Met Asn Phe Phe Gly Ser Leu Glu Ala  
100 105 110

Gln Lys Asn Leu Tyr Gly Asn Ser Tyr Ala Asn Ile Glu Phe Asp Arg  
115 120 125

Lys Gly Lys Val Gln Ala Leu Trp Pro Ile Asp Ala Ser Lys Val Thr  
130 135 140

Val Tyr Ile Asp Asp Val Gly Leu Leu Asn Ser Lys Thr Lys Met Trp  
145 150 155 160

Tyr Val Val Asn Thr Gly Gly Gln Gln Arg Val Leu Lys Pro Glu Glu  
165 170 175

Ile Leu His Phe Lys Asn Gly Ile Thr Leu Asp Gly Leu Val Gly Val  
180 185 190



PCT-US2005-009928\_Sequence Listing.txt as filed 10-3-05 in PCT

Pro Thr Met Glu Tyr Leu Lys Ser Thr Leu Glu Asn Ser Ala Ser Ala  
195 200 205

Asp Lys Phe Ile Asn Asn Phe Tyr Lys Gln Gly Leu Gln Val Lys Gly  
210 215 220

Leu Val Gln Tyr Val Gly Asp Leu Asn Glu Asp Ala Lys Lys Val Phe  
225 230 235 240

Arg Glu Asn Phe Glu Ser Met Ser Ser Gly Leu Gln Asn Ser His Arg  
245 250 255

Ile Ala Leu Met Pro Val Gly Tyr Gln Phe Gln Pro Ile Ser Leu Asn  
260 265 270

Met Ser Asp Ala Gln Phe Leu Glu Asn Thr Glu Leu Thr Ile Arg Gln  
275 280 285

Ile Ala Thr Ala Phe Gly Ile Lys Met His Gln Leu Asn Asp Leu Ser  
290 295 300

Lys Ala Thr Leu Asn Asn Ile Glu Gln Gln Gln Gln Gln Phe Tyr Thr  
305 310 315 320

Asp Thr Leu Gln Ala Thr Leu Thr Met Tyr Glu Gln Glu Met Thr Tyr  
325 330 335

Lys Leu Phe Leu Asp Ser Glu Leu Asp Lys Gly Phe Tyr Ser Lys Phe  
340 345 350

Asn Val Asp Ala Ile Leu Arg Ala Asp Ile Lys Thr Arg Tyr Glu Ala  
355 360 365

Tyr Arg Thr Gly Ile Gln Gly Gly Phe Leu Lys Pro Asn Glu Ala Arg  
370 375 380

Ser Lys Glu Asp Leu Pro Pro Glu Ala Gly Gly Asp Arg Leu Leu Val  
385 390 395 400

Asn Gly Asn Met Leu Pro Ile Asp Met Ala Gly Gln Ala Tyr Leu Lys  
405 410 415

Gly Gly Asp Thr Asn Gly Glu Val Ser Lys Glu Gly Asn Glu Gly Asn  
420 425 430

<210> 9  
<211> 206  
<212> PRT  
<213> Bacillus anthracis  
<400> 9

PCT-US2005-009928\_Sequence Listing.txt as filed 10-3-05 in PCT

Met Glu Lys Ser Ala Lys Lys Glu Met Lys Glu Ile Arg Ala Leu Pro  
1 5 10 15

Met Thr Ile Glu Val Arg Glu Val Asn Glu Asp Glu Gly Lys Arg Thr  
20 25 30

Ile Ser Gly Ser Ile Lys Tyr Asn Asn Glu Ser Ala Glu Met Arg Asp  
35 40 45

Trp Trp Gly Asp Thr Phe Val Glu Glu Ile Ala Glu Gly Ala Phe Asp  
50 55 60

Glu Ser Leu Lys Val Arg Asp Val Val Gly Leu Trp Ser His Asp Thr  
65 70 75 80

Ser Gln Val Leu Gly Asn Thr Lys Ser Lys Thr Leu Arg Ile Glu Asn  
85 90 95

Asp Lys Lys Glu Leu Arg Phe Glu Leu Asp Ile Pro Asn Thr Thr Val  
100 105 110

Gly Asn Asp Ala Trp Glu Leu Ile Lys Arg Gly Asp Val Asp Gly Val  
115 120 125

Ser Phe Gly Met Lys Val Thr Lys Asp Lys Trp Ser Ser Glu Glu Arg  
130 135 140

Glu Asn Gly Lys Leu Tyr Lys Arg Ser Ile Leu Asn Ala Glu Leu Tyr  
145 150 155 160

Glu Ile Ser Pro Val Ala Phe Pro Ala Tyr Pro Thr Asn Glu Val Ser  
165 170 175

Val Arg Ser Leu Asp Asp Phe Lys Ala Gly Glu Lys Arg Val Ala Asp  
180 185 190

Glu Phe Arg Lys Arg Lys Leu Gln Ile Glu Leu Glu Leu Ile  
195 200 205

<210> 10  
<211> 206  
<212> PRT  
<213> Bacillus anthracis

<400> 10

Met Glu Lys Ser Ala Lys Lys Glu Met Lys Glu Ile Arg Ala Leu Pro  
1 5 10 15

Met Thr Ile Glu Val Arg Glu Val Asn Glu Asp Glu Gly Lys Arg Thr  
20 25 30

Ile Ser Gly Ser Ile Lys Tyr Asn Asn Glu Ser Ala Glu Met Arg Asp  
35 40 45

Trp Trp Gly Asp Thr Phe Val Glu Glu Ile Ala Glu Gly Ala Phe Asp  
50 55 60

Glu Ser Leu Lys Val Arg Asp Val Val Gly Leu Trp Ser His Asp Thr  
65 70 75 80

Ser Gln Val Leu Gly Asn Thr Lys Ser Lys Thr Leu Arg Ile Glu Asn  
85 90 95

Asp Lys Lys Glu Leu Arg Phe Glu Leu Asp Ile Pro Asn Thr Thr Val  
100 105 110

Gly Asn Asp Ala Trp Glu Leu Ile Lys Arg Gly Asp Val Asp Gly Val  
115 120 125

Ser Phe Gly Met Lys Val Thr Lys Asp Lys Trp Ser Ser Glu Glu Arg  
130 135 140

Glu Asn Gly Lys Leu Tyr Lys Arg Ser Ile Leu Asn Ala Glu Leu Tyr  
145 150 155 160

Glu Ile Ser Pro Val Ala Phe Pro Ala Tyr Pro Thr Asn Glu Val Ser  
165 170 175

Val Arg Ser Leu Asp Asp Phe Lys Ala Gly Glu Lys Arg Val Ala Asp  
180 185 190

Glu Phe Arg Lys Arg Lys Leu Gln Ile Glu Leu Glu Leu Ile  
195 200 205

<210> 11  
<211> 392  
<212> PRT  
<213> Bacillus anthracis

<400> 11

Met Ser Lys Glu Leu Arg Glu Leu Leu Ala Lys Leu Glu Gly Lys Lys  
1 5 10 15

Glu Glu Val Arg Ser Leu Met Gly Glu Asp Lys Val Ala Glu Ala Glu  
20 25 30

Gln Met Met Glu Glu Val Arg Ser Leu Gln Lys Lys Ile Asp Leu Gln  
35 40 45

Arg Ser Leu Asp Glu Ala Glu Thr Glu Glu Arg Asn Asn Gly Arg Glu  
50 55 60

PCT-US2005-009928\_Sequence Listing.txt as filed 10-3-05 in PCT

Val	Glu	Thr	Arg	Asn	Val	Asp	Gly	Glu	Met	Glu	Tyr	Arg	Asp	Val	Phe
65					70					75					80
Met	Lys	Ala	Leu	Arg	Asn	Lys	Pro	Leu	Asn	Ala	Glu	Glu	Arg	Glu	Phe
				85					90					95	
Leu	Glu	Asp	Asp	Leu	Glu	Gln	Arg	Ala	Met	Ser	Gly	Leu	Thr	Gly	Glu
			100					105					110		
Asp	Gly	Gly	Leu	Val	Ile	Pro	Gln	Asp	Ile	Gln	Thr	Gln	Ile	Asn	Glu
		115					120					125			
Leu	Ala	Arg	Ser	Phe	Asp	Ala	Leu	Glu	Gln	Tyr	Val	Thr	Val	Glu	Pro
	130					135					140				
Val	Arg	Thr	Arg	Ser	Gly	Ser	Arg	Val	Leu	Glu	Lys	Asn	Ser	Asp	Met
145					150					155					160
Ile	Pro	Phe	Ala	Glu	Ile	Thr	Glu	Met	Gly	Glu	Ile	Pro	Glu	Thr	Asp
				165					170					175	
Asn	Pro	Lys	Phe	Ser	Asn	Val	Gln	Tyr	Ala	Val	Lys	Asp	Arg	Ala	Gly
			180					185					190		
Ile	Leu	Pro	Leu	Ser	Arg	Ser	Leu	Leu	Gln	Asp	Ser	Asp	Gln	Asn	Ile
		195					200					205			
Leu	Lys	Tyr	Val	Thr	Lys	Trp	Leu	Gly	Lys	Lys	Ser	Lys	Val	Thr	Arg
	210					215					220				
Asn	Val	Leu	Ile	Leu	Gly	Val	Ile	Glu	Lys	Leu	Thr	Lys	Gln	Ala	Ile
225					230					235					240
Lys	Ser	Leu	Asp	Asp	Ile	Lys	Asp	Val	Leu	Asn	Val	Lys	Leu	Asp	Pro
				245					250					255	
Ala	Ile	Ser	Pro	Asn	Ala	Ile	Leu	Leu	Thr	Asn	Gln	Asp	Gly	Phe	Asn
			260					265					270		
Tyr	Leu	Asp	Lys	Leu	Lys	Asp	Lys	Asp	Gly	Lys	Tyr	Ile	Leu	Gln	Ser
		275					280					285			
Asp	Pro	Thr	Gln	Lys	Asn	Lys	Lys	Leu	Phe	Ala	Gly	Thr	Asn	Pro	Val
	290					295					300				
Val	Val	Val	Ser	Asn	Arg	Phe	Leu	Lys	Ser	Lys	Gly	Thr	Thr	Ala	Lys
305					310					315					320
Lys	Ala	Pro	Leu	Ile	Ile	Gly	Asp	Leu	Lys	Glu	Ala	Ile	Val	Leu	Phe
				325					330					335	

Lys Arg Glu Asp Met Glu Leu Ala Ser Thr Asp Val Gly Gly Lys Ala  
340 345 350

Phe Thr Arg Asn Thr Leu Asp Leu Arg Ala Ile Gln Arg Asp Asp Val  
355 360 365

Gln Met Trp Asp Asn Glu Ala Ala Val Tyr Gly Glu Ile Asp Leu Ser  
370 375 380

Ala Pro Val Glu Gln Pro Gln Gly  
385 390

<210> 12  
<211> 392  
<212> PRT  
<213> Bacillus anthracis

<400> 12

Met Ser Lys Glu Leu Arg Glu Leu Leu Ala Lys Leu Glu Gly Lys Lys  
1 5 10 15

Glu Glu Val Arg Ser Leu Met Gly Glu Asp Lys Val Ala Glu Ala Glu  
20 25 30

Gln Met Met Glu Glu Val Arg Ser Leu Gln Lys Lys Ile Asp Leu Gln  
35 40 45

Arg Ser Leu Asp Glu Ala Glu Thr Glu Glu Arg Asn Asn Gly Arg Glu  
50 55 60

Val Glu Thr Arg Asn Val Asp Gly Glu Met Glu Tyr Arg Asp Val Phe  
65 70 75 80

Met Lys Ala Leu Arg Asn Lys Pro Leu Asn Ala Glu Glu Arg Glu Phe  
85 90 95

Leu Glu Asp Asp Leu Glu Gln Arg Ala Met Ser Gly Leu Thr Gly Glu  
100 105 110

Asp Gly Gly Leu Val Ile Pro Gln Asp Ile Gln Thr Gln Ile Asn Glu  
115 120 125

Leu Ala Arg Ser Phe Asp Ala Leu Glu Gln Tyr Val Thr Val Glu Pro  
130 135 140

Val Arg Thr Arg Ser Gly Ser Arg Val Leu Glu Lys Asn Ser Asp Met  
145 150 155 160

Ile Pro Phe Ala Glu Ile Thr Glu Met Gly Glu Ile Pro Glu Thr Asp  
165 170 175



PCT-US2005-009928\_Sequence Listing.txt as filed 10-3-05 in PCT

Asn Pro Lys Phe Ser Asn Val Gln Tyr Ala Val Lys Asp Arg Ala Gly  
180 185 190

Ile Leu Pro Leu Ser Arg Ser Leu Leu Gln Asp Ser Asp Gln Asn Ile  
195 200 205

Leu Lys Tyr Val Thr Lys Trp Leu Gly Lys Lys Ser Lys Val Thr Arg  
210 215 220

Asn Val Leu Ile Leu Gly Val Ile Glu Lys Leu Thr Lys Gln Ala Ile  
225 230 235 240

Lys Ser Leu Asp Asp Ile Lys Asp Val Leu Asn Val Lys Leu Asp Pro  
245 250 255

Ala Ile Ser Pro Asn Ala Ile Leu Leu Thr Asn Gln Asp Gly Phe Asn  
260 265 270

Tyr Leu Asp Lys Leu Lys Asp Lys Asp Gly Lys Tyr Ile Leu Gln Ser  
275 280 285

Asp Pro Thr Gln Lys Asn Lys Lys Leu Phe Ala Gly Thr Asn Pro Val  
290 295 300

Val Val Val Ser Asn Arg Phe Leu Lys Ser Lys Gly Thr Thr Ala Lys  
305 310 315 320

Lys Ala Pro Leu Ile Ile Gly Asp Leu Lys Glu Ala Ile Val Leu Phe  
325 330 335

Lys Arg Glu Asp Met Glu Leu Ala Ser Thr Asp Val Gly Gly Lys Ala  
340 345 350

Phe Thr Arg Asn Thr Leu Asp Leu Arg Ala Ile Gln Arg Asp Asp Val  
355 360 365

Gln Met Trp Asp Asn Glu Ala Ala Val Tyr Gly Glu Ile Asp Leu Ser  
370 375 380

Ala Pro Val Glu Gln Pro Gln Gly  
385 390

<210> 13  
<211> 96  
<212> PRT  
<213> Bacillus anthracis

<400> 13

Met Leu Val Thr Leu Glu Glu Ala Lys Glu Trp Ile Arg Val Asp Gly  
1 5 10 15

Asp Asp Asp Pro Thr Ile Thr Met Leu Ile Lys Ala Ala Glu Leu Tyr  
20 25 30

Ile Tyr Lys Ala Thr Gly Lys Thr Phe Thr Gln Thr Asn Glu Asp Ala  
35 40 45

Lys Leu Leu Cys Leu Phe Leu Val Ala Asp Trp Tyr Gly Asn Arg Leu  
50 55 60

Leu Val Gly Glu Lys Ala Ser Glu Lys Ile Arg Thr Ile Val Gln Ser  
65 70 75 80

Met Ile Leu Gln Leu Gln Tyr Ala Ser Glu Pro Gln Glu Glu Arg Lys  
85 90 95

<210> 14  
<211> 96  
<212> PRT  
<213> Bacillus anthracis

<400> 14

Met Leu Val Thr Leu Glu Glu Ala Lys Glu Trp Ile Arg Val Asp Gly  
1 5 10 15

Asp Asp Asp Pro Thr Ile Thr Met Leu Ile Lys Ala Ala Glu Leu Tyr  
20 25 30

Ile Tyr Lys Ala Thr Gly Lys Thr Phe Thr Gln Thr Asn Glu Asp Ala  
35 40 45

Lys Leu Leu Cys Leu Phe Leu Val Ala Asp Trp Tyr Gly Asn Arg Leu  
50 55 60

Leu Val Gly Glu Lys Ala Ser Glu Lys Ile Arg Thr Ile Val Gln Ser  
65 70 75 80

Met Ile Leu Gln Leu Gln Tyr Ala Ser Glu Pro Gln Glu Glu Arg Lys  
85 90 95

<210> 15  
<211> 107  
<212> PRT  
<213> Bacillus anthracis

<400> 15

Met Asn Pro Ala Lys Leu Asp Lys Arg Leu Thr Phe Gln Val Lys Asp  
1 5 10 15

Glu Asn Ala Lys Gly Pro Asp Gly Asp Pro Ile Asp Gly Tyr Lys Asp  
20 25 30

Ala Phe Thr Val Trp Gly Ser Phe Val Tyr Leu Lys Gly Arg Lys Tyr  
35 40 45

Phe Glu Ala Ala Ala Ala Asn Ser Glu Val Gln Gly Glu Thr Glu Ile  
50 55 60

Arg Asn Arg Asp Asp Val Ser Ala Asp Met Lys Ile Lys Tyr Lys Asn  
65 70 75 80

Val Ile Tyr Asp Ile Val Ser Val Ile Pro Thr Gln Asp His Thr Leu  
85 90 95

Leu Ile Met Trp Lys Arg Gly Glu Met Asn Gly  
100 105

<210> 16  
<211> 107  
<212> PRT  
<213> Bacillus anthracis

<400> 16

Met Asn Pro Ala Lys Leu Asp Lys Arg Leu Thr Phe Gln Val Lys Asp  
1 5 10 15

Glu Asn Ala Lys Gly Pro Asp Gly Asp Pro Ile Asp Gly Tyr Lys Asp  
20 25 30

Ala Phe Thr Val Trp Gly Ser Phe Val Tyr Leu Lys Gly Arg Lys Tyr  
35 40 45

Phe Glu Ala Ala Ala Ala Asn Ser Glu Val Gln Gly Glu Thr Glu Ile  
50 55 60

Arg Asn Arg Asp Asp Val Ser Ala Asp Met Lys Ile Lys Tyr Lys Asn  
65 70 75 80

Val Ile Tyr Asp Ile Val Ser Val Ile Pro Thr Gln Asp His Thr Leu  
85 90 95

Leu Ile Met Trp Lys Arg Gly Glu Met Asn Gly  
100 105

<210> 17  
<211> 105  
<212> PRT  
<213> Bacillus anthracis

<400> 17

Met Lys Leu Thr Leu Met Ile Asn Lys Glu Lys Gln Thr Phe Asn Met  
1 5 10 15

PCT-US2005-009928\_Sequence Listing.txt as filed 10-3-05 in PCT  
Pro Glu Phe Ile Pro Ala Arg Leu Ile Arg Gln Ala Pro Glu Leu Ala  
20 25 30

Glu Ile Pro Asn Asn Pro Gly Pro Glu Asp Met Asp Lys Met Val Gln  
35 40 45

Phe Val Val Lys Val Tyr Asp Gly Gln Phe Thr Leu Asp Gln Tyr Trp  
50 55 60

Asp Gly Val Asp Ala Arg Lys Phe Leu Ser Thr Thr Ser Asp Val Ile  
65 70 75 80

Asn Ala Ile Ile Asn Glu Thr Val Glu Ala Ala Gly Gly Ser Thr Glu  
85 90 95

Ser Gly Glu Glu Glu Asn Pro Asn Ala  
100 105

<210> 18  
<211> 105  
<212> PRT  
<213> Bacillus anthracis

<400> 18

Met Lys Leu Thr Leu Met Ile Asn Lys Glu Lys Gln Thr Phe Asn Met  
1 5 10 15

Pro Glu Phe Ile Pro Ala Arg Leu Ile Arg Gln Ala Pro Glu Leu Ala  
20 25 30

Glu Ile Pro Asn Asn Pro Gly Pro Glu Asp Met Asp Lys Met Val Gln  
35 40 45

Phe Val Val Lys Val Tyr Asp Gly Gln Phe Thr Leu Asp Gln Tyr Trp  
50 55 60

Asp Gly Val Asp Ala Arg Lys Phe Leu Ser Thr Thr Ser Asp Val Ile  
65 70 75 80

Asn Ala Ile Ile Asn Glu Thr Val Glu Ala Ala Gly Gly Ser Thr Glu  
85 90 95

Ser Gly Glu Glu Glu Asn Pro Asn Ala  
100 105

<210> 19  
<211> 119  
<212> PRT  
<213> Bacillus anthracis

<400> 19

Val Ile Asn Leu Arg Pro Asp Ile Leu Gln Ala Leu Glu Asn Asp Gln  
Page 57

Glu Leu Val Ser Leu Leu Gly Gly Lys Arg Ile Tyr Tyr Arg Lys Ala  
20 25 30

Lys Lys Ala Glu Glu Phe Pro Arg Ile Thr Tyr Phe Glu Leu Asp Asn  
35 40 45

Arg Pro Asp Gly Phe Ala Asp Asn Gln Glu Ile Glu Ser Glu Ile Leu  
50 55 60

Phe Gln Val Asp Val Trp Ala Lys Ser Ser Thr Thr Ala Ile His Gln  
65 70 75 80

Lys Val Asn Glu Ile Met Lys Arg Ile Gly Phe Ser Arg Tyr Ala Val  
85 90 95

Ala Asp Leu Tyr Glu Glu Asp Thr Gln Ile Phe His Tyr Ala Met Arg  
100 105 110

Phe Ala Lys Gly Val Glu Leu  
115

<210> 20  
<211> 119  
<212> PRT  
<213> Bacillus anthracis

<400> 20

Val Ile Asn Leu Arg Pro Asp Ile Leu Gln Ala Leu Glu Asn Asp Gln  
1 5 10 15

Glu Leu Val Ser Leu Leu Gly Gly Lys Arg Ile Tyr Tyr Arg Lys Ala  
20 25 30

Lys Lys Ala Glu Glu Phe Pro Arg Ile Thr Tyr Phe Glu Leu Asp Asn  
35 40 45

Arg Pro Asp Gly Phe Ala Asp Asn Gln Glu Ile Glu Ser Glu Ile Leu  
50 55 60

Phe Gln Val Asp Val Trp Ala Lys Ser Ser Thr Thr Ala Ile His Gln  
65 70 75 80

Lys Val Asn Glu Ile Met Lys Arg Ile Gly Phe Ser Arg Tyr Ala Val  
85 90 95

Ala Asp Leu Tyr Glu Glu Asp Thr Gln Ile Phe His Tyr Ala Met Arg  
100 105 110

Phe Ala Lys Gly Val Glu Leu



<210> 21  
<211> 202  
<212> PRT  
<213> Bacillus anthracis

<400> 21

Met Ala Gly Glu Val Val Arg Ile Ser Ser Thr Val Gly Val Asp Asn  
1 5 10 15

Leu Val Tyr Ala Lys Val Leu Gln Asp Asp Ser Ser Ala Ile Lys Tyr  
20 25 30

Thr Asp Val Lys Lys Met Glu Gly Ala Val Lys Val Lys Leu Thr Lys  
35 40 45

Lys Val Ala Ser Glu Val Met Trp Ser Asp Asn Arg Lys Ser Glu Ile  
50 55 60

Ala Glu Ser Asp Gly Glu Thr Glu Val Glu Ile Glu Val Arg Gly Leu  
65 70 75 80

Ser Leu Ser Thr Lys Ala Asp Ile Glu Gly Phe Pro Glu Val Lys Asp  
85 90 95

Gly Val Leu Asp Glu Lys Arg Glu Gly Glu Lys Pro Tyr Leu Ala Ile  
100 105 110

Gly Phe Arg Phe Leu Lys Ala Asn Asp Lys Tyr Arg Tyr Val Trp Leu  
115 120 125

Leu Lys Gly Lys Leu Ser Gln Glu Glu Glu Glu Ala Glu Thr Lys Lys  
130 135 140

Asp Lys Pro Asn Phe Gln Thr Thr Lys Leu Lys Gly Ser Phe Ile Glu  
145 150 155 160

Arg Asp Phe Asp Asp Arg Thr Lys Phe Thr Ala Asp Glu Asp Glu Pro  
165 170 175

Thr Phe Thr Lys Leu Val Gly Asp Asn Trp Phe Asn Lys Val Tyr Glu  
180 185 190

Lys Pro Val Thr Gln Pro Pro Ala Gly Lys  
195 200

<210> 22  
<211> 202  
<212> PRT  
<213> Bacillus anthracis

<400> 22

Met Ala Gly Glu Val Val Arg Ile Ser Ser Thr Val Gly Val Asp Asn  
1 5 10 15

Leu Val Tyr Ala Lys Val Leu Gln Asp Asp Ser Ser Ala Ile Lys Tyr  
20 25 30

Thr Asp Val Lys Lys Met Glu Gly Ala Val Lys Val Lys Leu Thr Lys  
35 40 45

Lys Val Ala Ser Glu Val Met Trp Ser Asp Asn Arg Lys Ser Glu Ile  
50 55 60

Ala Glu Ser Asp Gly Glu Thr Glu Val Glu Ile Glu Val Arg Gly Leu  
65 70 75 80

Ser Leu Ser Thr Lys Ala Asp Ile Glu Gly Phe Pro Glu Val Lys Asp  
85 90 95

Gly Val Leu Asp Glu Lys Arg Glu Gly Glu Lys Pro Tyr Leu Ala Ile  
100 105 110

Gly Phe Arg Phe Leu Lys Ala Asn Asp Lys Tyr Arg Tyr Val Trp Leu  
115 120 125

Leu Lys Gly Lys Leu Ser Gln Glu Glu Glu Glu Ala Glu Thr Lys Lys  
130 135 140

Asp Lys Pro Asn Phe Gln Thr Thr Lys Leu Lys Gly Ser Phe Ile Glu  
145 150 155 160

Arg Asp Phe Asp Asp Arg Thr Lys Phe Thr Ala Asp Glu Asp Glu Pro  
165 170 175

Thr Phe Thr Lys Leu Val Gly Asp Asn Trp Phe Asn Lys Val Tyr Glu  
180 185 190

Lys Pro Val Thr Gln Pro Pro Ala Gly Lys  
195 200

<210> 23

<211> 105

<212> PRT

<213> Bacillus anthracis

<400> 23

Met Lys Leu Thr Leu Met Ile Asn Lys Glu Lys Gln Thr Phe Asn Met  
1 5 10 15

Pro Glu Phe Ile Pro Ala Arg Leu Ile Arg Gln Ala Pro Glu Leu Ala  
20 25 30

Glu Ile Pro Asn Asn Pro Gly Pro Glu Asp Met Asp Lys Met Val Gln  
35 40 45

Phe Val Val Lys Val Tyr Asp Gly Gln Phe Thr Leu Asp Gln Tyr Trp  
50 55 60

Asp Gly Val Asp Ala Arg Lys Phe Leu Ser Thr Thr Ser Asp Val Ile  
65 70 75 80

Asn Ala Ile Ile Asn Glu Thr Val Glu Ala Ala Gly Gly Ser Thr Glu  
85 90 95

Ser Gly Glu Glu Glu Asn Pro Asn Ala  
100 105

<210> 24  
<211> 105  
<212> PRT  
<213> Bacillus anthracis

<400> 24

Met Lys Leu Thr Leu Met Ile Asn Lys Glu Lys Gln Thr Phe Asn Met  
1 5 10 15

Pro Glu Phe Ile Pro Ala Arg Leu Ile Arg Gln Ala Pro Glu Leu Ala  
20 25 30

Glu Ile Pro Asn Asn Pro Gly Pro Glu Asp Met Asp Lys Met Val Gln  
35 40 45

Phe Val Val Lys Val Tyr Asp Gly Gln Phe Thr Leu Asp Gln Tyr Trp  
50 55 60

Asp Gly Val Asp Ala Arg Lys Phe Leu Ser Thr Thr Ser Asp Val Ile  
65 70 75 80

Asn Ala Ile Ile Asn Glu Thr Val Glu Ala Ala Gly Gly Ser Thr Glu  
85 90 95

Ser Gly Glu Glu Glu Asn Pro Asn Ala  
100 105

<210> 25  
<211> 58  
<212> PRT  
<213> Bacillus anthracis

<400> 25

Met Asp Glu Leu Tyr Leu Ser Leu Leu Arg Gln Gly Tyr Lys His His  
1 5 10 15

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His Ile Asp Asn Glu Met Asp Ile Trp His Tyr Leu Arg Leu Asn Arg  
20 25 30

Lys Met His Glu Asn Gly Asn Glu Asn Tyr Glu Gly Ser Asn Ser Asn  
35 40 45

Glu Ile Glu Val Pro Ala Glu Asn Ile Ile  
50 55

<210> 26  
<211> 58  
<212> PRT  
<213> Bacillus anthracis

<400> 26

Met Asp Glu Leu Tyr Leu Ser Leu Leu Arg Gln Gly Tyr Lys His His  
1 5 10 15

His Ile Asp Asn Glu Met Asp Ile Trp His Tyr Leu Arg Leu Asn Arg  
20 25 30

Lys Met His Glu Asn Gly Asn Glu Asn Tyr Glu Gly Ser Asn Ser Asn  
35 40 45

Glu Ile Glu Val Pro Ala Glu Asn Ile Ile  
50 55

<210> 27  
<211> 1283  
<212> PRT  
<213> Bacillus anthracis

<400> 27

Met Ala Asn Glu Ile Asn Asn Leu Val Val Arg Leu Ser Leu Asp Asn  
1 5 10 15

Val Asn Phe Arg Gln Gly Ile Ser Asn Ser Gly Arg Ala Val Arg Thr  
20 25 30

Leu Gln Asn Glu Leu Lys Ser Val Ser Thr Gly Met Gly Gly Phe Ala  
35 40 45

Asn Ala Ser Gln Gln Thr Gln Ala Lys Met Asn Thr Leu Ser Arg Leu  
50 55 60

Ile Asp Ala Gln Lys Glu Lys Val Lys Ala Leu Arg Gln Ala Tyr Asp  
65 70 75 80

Gln Asn Lys Ala Lys Leu Gly Glu Asn Asp Ala Ala Thr Gln Arg Tyr  
85 90 95

PCT-US2005-009928\_Sequence Listing.txt as filed 10-3-05 in PCT  
Ala Ser Gln Val Asn Lys Ala Val Ala Asp Leu Asn Arg Phe Glu Asn  
100 105 110

Glu Leu Lys Gln Val Asn Arg Gln Ala Glu Gln Lys Gly Met Asp Lys  
115 120 125

Leu Asn Asn Ser Leu Lys Ser Leu Gln Ala Glu Phe Gln Ser Ile Thr  
130 135 140

Thr Gly Met Gly Gly Phe Ser Asn Ala Thr Glu Gln Thr Arg Ala Lys  
145 150 155 160

Val Asp Val Leu Ser Arg Met Val Asp Lys Gln Lys Glu Lys Ile Arg  
165 170 175

Glu Leu Gln Gln Ala Tyr Asn Arg Ala Lys Thr Glu Glu Gly Glu Ala  
180 185 190

Ser Gln Ser Ala Gln Arg Tyr Ala Glu Gln Ile His Arg Ala Thr Ala  
195 200 205

Glu Leu Asn Arg Phe Glu Thr Gly Leu Gln Gln Ser Asn Arg Glu Leu  
210 215 220

Glu Gln Gln Gly Asn Arg Leu Leu Asn Phe Gly Asn Arg Met Glu Thr  
225 230 235 240

Leu Gly Asn His Leu Gln Asn Ala Gly Met Gln Ile Gly Met Val Phe  
245 250 255

Gly Gly Met Thr Tyr Ala Ile Gly Arg Gly Leu Lys Ser Ala Ile Thr  
260 265 270

Glu Ser Met Asn Phe Glu Gln Gln Met Ala Asn Val Lys Ala Val Ser  
275 280 285

Gly Ser Thr Gly Ala Glu Met Lys Lys Leu Ser Glu Leu Ala Val Asn  
290 295 300

Met Gly Glu Thr Thr Lys Tyr Ser Ser Val Gln Ala Gly Gln Gly Ile  
305 310 315 320

Glu Glu Leu Ile Lys Ala Gly Val Ser Leu Gln Asp Ile Ile Asn Gly  
325 330 335

Gly Leu Ala Gly Ala Leu Asn Leu Ala Thr Ala Gly Glu Leu Glu Leu  
340 345 350

Gly Glu Ala Ala Glu Ile Ala Ser Thr Ala Leu Asn Ala Phe Lys Ala  
355 360 365



Asp His Leu Ser Val Ala Asp Ala Ala Asn Ile Leu Ser Gly Ala Ala  
 370 375 380  
 Asn Ala Ser Ala Thr Asp Val Arg Glu Leu Lys Tyr Gly Leu Ser Ala  
 385 390 395 400  
 Ser Ser Ala Val Ala Ala Gly Ala Gly Met Thr Phe Lys Asp Thr Ala  
 405 410 415  
 Thr Thr Leu Ala Val Phe Ala Gln Asn Gly Leu Lys Gly Ser Asp Ala  
 420 425 430  
 Gly Thr Ser Leu Lys Thr Met Leu Met Arg Leu Asn Pro Ser Thr Lys  
 435 440 445  
 Glu Ala Tyr Asn Lys Met Arg Asp Leu Gly Leu Ile Thr Tyr Asn Ala  
 450 455 460  
 Gln Ala Gly Phe Asp Phe Leu Val Lys Asn Gly Ile Gln Pro Ala Ser  
 465 470 475 480  
 Arg Asn Val Gly Asp Ile Glu Val Ala Leu Glu Gln Tyr Val Met Lys  
 485 490 495  
 Thr Glu Gly Val Thr Lys Trp Asn Asp Lys Cys Asp Thr Thr Phe Arg  
 500 505 510  
 Glu Leu Ala Thr Ser Ser Ala Phe Leu Ser Ser Lys Phe Tyr Asp Gln  
 515 520 525  
 Gln Gly His Ile Gln Ser Leu Glu Asn Ile Ser Gly Thr Leu His Glu  
 530 535 540  
 Ser Met Lys Asp Leu Thr Asp Gln Gln Arg Ser Met Ala Leu Glu Thr  
 545 550 555 560  
 Leu Phe Gly Ser Asp Ala Val Arg Gly Ala Thr Ile Leu Phe Lys Glu  
 565 570 575  
 Gly Ala Lys Gly Val Asn Glu Met Trp Asp Ser Met Ser Lys Val Thr  
 580 585 590  
 Ala Ala Asp Val Ala Ala Thr Lys Ile Asp Thr Leu Lys Gly Arg Leu  
 595 600 605  
 Thr Leu Leu Asp Ser Ala Phe Ser Thr Met Lys Lys Thr Ile Gly Asp  
 610 615 620  
 Ala Leu Ala Pro Val Val Ser Val Phe Val Ala Gly Leu Gln Lys Leu  
 625 630 635 640

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Val	Asp	Gly	Phe	Asn	Ser	Leu	Pro	Gly	Pro	Val	Gln	Lys	Ala	Ile	Ala		
				645					650					655			
Ile	Thr	Gly	Gly	Ile	Val	Leu	Ala	Leu	Thr	Ala	Val	Ala	Thr	Ala	Ile		
			660					665					670				
Gly	Val	Val	Leu	Ala	Ala	Phe	Gly	Met	Ile	Ala	Ser	Gly	Ile	Gly	Ser		
		675					680					685					
Leu	Ser	Leu	Ala	Leu	Ala	Ser	Val	Gly	Gly	Ile	Ala	Gly	Ile	Ala	Ala		
	690					695						700					
Gly	Ala	Val	Gly	Phe	Leu	Gly	Ser	Ala	Leu	Ala	Val	Leu	Thr	Gly	Pro		
705					710					715					720		
Ile	Gly	Leu	Val	Ala	Ala	Ala	Leu	Ile	Gly	Thr	Gly	Val	Val	Ala	Tyr		
				725					730					735			
Lys	Ala	Tyr	Gln	Lys	Ala	Thr	Glu	Asp	Ser	Ile	Ala	Ser	Val	Asp	Arg		
			740					745					750				
Phe	Ala	Thr	Asn	Thr	Glu	Gly	Lys	Val	Ser	Ser	Ser	Thr	Lys	Lys	Val		
		755					760					765					
Leu	Gly	Glu	Tyr	Phe	Lys	Leu	Ser	Asp	Gly	Ile	Arg	Gln	Lys	Leu	Thr		
	770					775					780						
Glu	Ile	Arg	Leu	Asn	His	Glu	Val	Ile	Thr	Glu	Glu	Gln	Ser	Gln	Lys		
785					790					795					800		
Leu	Ile	Gly	Gln	Tyr	Asp	Lys	Leu	Ala	Asn	Thr	Ile	Ile	Glu	Lys	Thr		
				805					810					815			
Asn	Ala	Arg	Gln	Gln	Lys	Glu	Ile	Glu	Gly	Leu	Lys	Lys	Phe	Phe	Ala		
			820					825					830				
Asp	Ser	Tyr	Val	Leu	Thr	Ala	Glu	Glu	Glu	Asn	Lys	Arg	Ile	Glu	Gln		
		835					840					845					
Leu	Asn	Gln	His	Tyr	Glu	Gln	Glu	Lys	Leu	Lys	Thr	Gln	Glu	Lys	Glu		
	850					855					860						
Asn	Lys	Ile	Lys	Glu	Ile	Leu	Gln	Thr	Ala	Ala	Arg	Glu	Asn	Arg	Glu		
865					870					875					880		
Leu	Thr	Thr	Ser	Glu	Arg	Ile	Ser	Leu	Gln	Ala	Leu	Gln	Asp	Glu	Met		
				885					890					895			
Asp	Arg	Val	Ala	Val	Glu	His	Met	Ser	Lys	Asn	Gln	Met	Glu	Gln	Lys		

Val Ile Leu Glu Asn Met Arg Val Gln Ala Ser Glu Ile Ser Ala Arg  
915 920 925

Gln Ala Ala Glu Val Val Glu Asn Ser Ala Lys Ala Arg Asp Lys Val  
930 935 940

Ile Glu Asp Ala Lys Lys Thr Arg Asp Glu Lys Ile Ala Glu Ala Ile  
945 950 955 960

Arg Gln Arg Asp Glu Asn Lys Thr Ile Thr Ala Asp Glu Ala Asn Ala  
965 970 975

Ile Ile Ala Glu Ala Lys Arg Gln Tyr Asp Ser Thr Val Ser Thr Ala  
980 985 990

Arg Asp Lys His Lys Glu Ile Val Ser Glu Ala Lys Ala Gln Ala Gly  
995 1000 1005

Glu His Ala Asn Gln Val Asp Trp Glu Thr Gly Gln Val Lys Ser  
1010 1015 1020

Lys Tyr Gln Ala Met Lys Asp Asp Val Ile Arg Lys Met Lys Glu  
1025 1030 1035

Met Trp Ser Asp Val Thr Asn Lys Tyr Glu Asp Met Lys Asn Ser  
1040 1045 1050

Ala Ser Asn Lys Val Glu Glu Ile Lys Asn Thr Val Ser Arg Lys  
1055 1060 1065

Phe Glu Glu Gln Lys Lys Ala Val Thr Asp Lys Met Ser Glu Ile  
1070 1075 1080

Lys Ser Ser Ile Glu Asp Lys Trp Asn Thr Val Glu Lys Phe Phe  
1085 1090 1095

Ser Ser Ile Asn Leu Arg Ser Ile Gly Lys Ser Ile Ile Glu Gly  
1100 1105 1110

Leu Gly Lys Gly Ile Asp Asp Ala Ser Gly Gly Leu Phe Ser Lys  
1115 1120 1125

Ala Ala Glu Ile Ala Ser Asp Ile Lys Lys Thr Ile Ser Gly Ala  
1130 1135 1140

Leu Glu Ile Asn Ser Pro Ser Lys Val Met Ile Pro Val Gly Ser  
1145 1150 1155

Ala Val Pro Glu Gly Val Gly Val Gly Met Asp Lys Gly Lys Arg  
1160 1165 1170

Phe Val Val Asp Ala Ala Lys Asn Val Val Gly Thr Val Lys Lys  
1175 1180 1185

Gln Met Gly Asn Met Pro Ser Val Phe Asp Phe Gly Phe Gln Thr  
1190 1195 1200

Asn Gln Tyr Ser Ile Pro Gln Asn Thr Phe Ser Asp Phe Ser Gly  
1205 1210 1215

Tyr Met Gln Pro Gln Leu Ser Tyr Asn Asn Pro Ser Met Ala Lys  
1220 1225 1230

Thr Ile Phe Pro Asn Arg Pro Gly Gly Glu Gln Glu Leu Asn Leu  
1235 1240 1245

Thr Val Asn Met Thr Asn Val Leu Asp Gly Lys Glu Leu Ala Asn  
1250 1255 1260

Gly Ser Tyr Thr Tyr Thr Thr Lys Leu Gln Asn Arg Glu Gln Lys  
1265 1270 1275

Arg Arg Ala Glu Phe  
1280

<210> 28  
<211> 1283  
<212> PRT  
<213> Bacillus anthracis

<400> 28

Met Ala Asn Glu Ile Asn Asn Leu Val Val Arg Leu Ser Leu Asp Asn  
1 5 10 15

Val Asn Phe Arg Gln Gly Ile Ser Asn Ser Gly Arg Ala Val Arg Thr  
20 25 30

Leu Gln Asn Glu Leu Lys Ser Val Ser Thr Gly Met Gly Gly Phe Ala  
35 40 45

Asn Ala Ser Gln Gln Thr Gln Ala Lys Met Asn Thr Leu Ser Arg Leu  
50 55 60

Ile Asp Ala Gln Lys Glu Lys Val Lys Ala Leu Arg Gln Ala Tyr Asp  
65 70 75 80

Gln Asn Lys Ala Lys Leu Gly Glu Asn Asp Ala Ala Thr Gln Arg Tyr  
85 90 95





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Asp His Leu Ser Val Ala Asp Ala Ala Asn Ile Leu Ser Gly Ala Ala  
 370 375 380  
 Asn Ala Ser Ala Thr Asp Val Arg Glu Leu Lys Tyr Gly Leu Ser Ala  
 385 390 395 400  
 Ser Ser Ala Val Ala Ala Gly Ala Gly Met Thr Phe Lys Asp Thr Ala  
 405 410 415  
 Thr Thr Leu Ala Val Phe Ala Gln Asn Gly Leu Lys Gly Ser Asp Ala  
 420 425 430  
 Gly Thr Ser Leu Lys Thr Met Leu Met Arg Leu Asn Pro Ser Thr Lys  
 435 440 445  
 Glu Ala Tyr Asn Lys Met Arg Asp Leu Gly Leu Ile Thr Tyr Asn Ala  
 450 455 460  
 Gln Ala Gly Phe Asp Phe Leu Val Lys Asn Gly Ile Gln Pro Ala Ser  
 465 470 475 480  
 Arg Asn Val Gly Asp Ile Glu Val Ala Leu Glu Gln Tyr Val Met Lys  
 485 490 495  
 Thr Glu Gly Val Thr Lys Trp Asn Asp Lys Cys Asp Thr Thr Phe Arg  
 500 505 510  
 Glu Leu Ala Thr Ser Ser Ala Phe Leu Ser Ser Lys Phe Tyr Asp Gln  
 515 520 525  
 Gln Gly His Ile Gln Ser Leu Glu Asn Ile Ser Gly Thr Leu His Glu  
 530 535 540  
 Ser Met Lys Asp Leu Thr Asp Gln Gln Arg Ser Met Ala Leu Glu Thr  
 545 550 555 560  
 Leu Phe Gly Ser Asp Ala Val Arg Gly Ala Thr Ile Leu Phe Lys Glu  
 565 570 575  
 Gly Ala Lys Gly Val Asn Glu Met Trp Asp Ser Met Ser Lys Val Thr  
 580 585 590  
 Ala Ala Asp Val Ala Ala Thr Lys Ile Asp Thr Leu Lys Gly Arg Leu  
 595 600 605  
 Thr Leu Leu Asp Ser Ala Phe Ser Thr Met Lys Lys Thr Ile Gly Asp  
 610 615 620  
 Ala Leu Ala Pro Val Val Ser Val Phe Val Ala Gly Leu Gln Lys Leu  
 625 630 635 640

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Val	Asp	Gly	Phe	Asn 645	Ser	Leu	Pro	Gly	Pro 650	Val	Gln	Lys	Ala	Ile 655	Ala
Ile	Thr	Gly	Gly 660	Ile	Val	Leu	Ala	Leu 665	Thr	Ala	Val	Ala	Thr 670	Ala	Ile
Gly	Val	Val 675	Leu	Ala	Ala	Phe	Gly 680	Met	Ile	Ala	Ser	Gly 685	Ile	Gly	Ser
Leu	Ser 690	Leu	Ala	Leu	Ala	Ser 695	Val	Gly	Gly	Ile	Ala 700	Gly	Ile	Ala	Ala
Gly 705	Ala	Val	Gly	Phe	Leu 710	Gly	Ser	Ala	Leu	Ala 715	Val	Leu	Thr	Gly	Pro 720
Ile	Gly	Leu	Val	Ala 725	Ala	Ala	Leu	Ile	Gly 730	Thr	Gly	Val	Val	Ala 735	Tyr
Lys	Ala	Tyr	Gln 740	Lys	Ala	Thr	Glu	Asp 745	Ser	Ile	Ala	Ser	Val 750	Asp	Arg
Phe	Ala	Thr 755	Asn	Thr	Glu	Gly	Lys 760	Val	Ser	Ser	Ser	Thr 765	Lys	Lys	Val
Leu	Gly 770	Glu	Tyr	Phe	Lys	Leu 775	Ser	Asp	Gly	Ile	Arg 780	Gln	Lys	Leu	Thr
Glu 785	Ile	Arg	Leu	Asn 790	His	Glu	Val	Ile	Thr	Glu 795	Glu	Gln	Ser	Gln	Lys 800
Leu	Ile	Gly	Gln 805	Tyr	Asp	Lys	Leu	Ala	Asn 810	Thr	Ile	Ile	Glu	Lys 815	Thr
Asn	Ala	Arg	Gln 820	Gln	Lys	Glu	Ile	Glu 825	Gly	Leu	Lys	Lys	Phe 830	Phe	Ala
Asp	Ser	Tyr 835	Val	Leu	Thr	Ala	Glu 840	Glu	Glu	Asn	Lys	Arg 845	Ile	Glu	Gln
Leu	Asn 850	Gln	His	Tyr	Glu	Gln 855	Glu	Lys	Leu	Lys	Thr 860	Gln	Glu	Lys	Glu
Asn 865	Lys	Ile	Lys	Glu	Ile 870	Leu	Gln	Thr	Ala	Ala 875	Arg	Glu	Asn	Arg	Glu 880
Leu	Thr	Thr	Ser	Glu 885	Arg	Ile	Ser	Leu	Gln 890	Ala	Leu	Gln	Asp	Glu	Met 895
Asp	Arg	Val	Ala	Val	Glu	His	Met	Ser	Lys	Asn	Gln	Met	Glu	Gln	Lys

Val Ile Leu Glu Asn Met Arg Val Gln Ala Ser Glu Ile Ser Ala Arg  
915 920 925

Gln Ala Ala Glu Val Val Glu Asn Ser Ala Lys Ala Arg Asp Lys Val  
930 935 940

Ile Glu Asp Ala Lys Lys Thr Arg Asp Glu Lys Ile Ala Glu Ala Ile  
945 950 955 960

Arg Gln Arg Asp Glu Asn Lys Thr Ile Thr Ala Asp Glu Ala Asn Ala  
965 970 975

Ile Ile Ala Glu Ala Lys Arg Gln Tyr Asp Ser Thr Val Ser Thr Ala  
980 985 990

Arg Asp Lys His Lys Glu Ile Val Ser Glu Ala Lys Ala Gln Ala Gly  
995 1000 1005

Glu His Ala Asn Gln Val Asp Trp Glu Thr Gly Gln Val Lys Ser  
1010 1015 1020

Lys Tyr Gln Ala Met Lys Asp Asp Val Ile Arg Lys Met Lys Glu  
1025 1030 1035

Met Trp Ser Asp Val Thr Asn Lys Tyr Glu Asp Met Lys Asn Ser  
1040 1045 1050

Ala Ser Asn Lys Val Glu Glu Ile Lys Asn Thr Val Ser Arg Lys  
1055 1060 1065

Phe Glu Glu Gln Lys Lys Ala Val Thr Asp Lys Met Ser Glu Ile  
1070 1075 1080

Lys Ser Ser Ile Glu Asp Lys Trp Asn Thr Val Glu Lys Phe Phe  
1085 1090 1095

Ser Ser Ile Asn Leu Arg Ser Ile Gly Lys Ser Ile Ile Glu Gly  
1100 1105 1110

Leu Gly Lys Gly Ile Asp Asp Ala Ser Gly Gly Leu Phe Ser Lys  
1115 1120 1125

Ala Ala Glu Ile Ala Ser Asp Ile Lys Lys Thr Ile Ser Gly Ala  
1130 1135 1140

Leu Glu Ile Asn Ser Pro Ser Lys Val Met Ile Pro Val Gly Ser  
1145 1150 1155

PCT-US2005-009928\_Sequence Listing.txt as filed 10-3-05 in PCT  
Ala Val Pro Glu Gly Val Gly Val Gly Met Asp Lys Gly Lys Arg  
1160 1165 1170

Phe Val Val Asp Ala Ala Lys Asn Val Val Gly Thr Val Lys Lys  
1175 1180 1185

Gln Met Gly Asn Met Pro Ser Val Phe Asp Phe Gly Phe Gln Thr  
1190 1195 1200

Asn Gln Tyr Ser Ile Pro Gln Asn Thr Phe Ser Asp Phe Ser Gly  
1205 1210 1215

Tyr Met Gln Pro Gln Leu Ser Tyr Asn Asn Pro Ser Met Ala Lys  
1220 1225 1230

Thr Ile Phe Pro Asn Arg Pro Gly Gly Glu Gln Glu Leu Asn Leu  
1235 1240 1245

Thr Val Asn Met Thr Asn Val Leu Asp Gly Lys Glu Leu Ala Asn  
1250 1255 1260

Gly Ser Tyr Thr Tyr Thr Thr Lys Leu Gln Asn Arg Glu Gln Lys  
1265 1270 1275

Arg Arg Ala Glu Phe  
1280

<210> 29  
<211> 496  
<212> PRT  
<213> Bacillus anthracis

<400> 29

Leu Gly Lys Leu Ser Phe Thr Phe Asn Asn Ile Arg Lys Asp Tyr Ile  
1 5 10 15

Gln Met Leu Val Gly Arg Lys Arg Pro Ser Trp Ala Pro Val Lys Arg  
20 25 30

Arg Leu Val Arg Val Pro His Arg Ala Gly Ala Leu Leu Leu Asn Thr  
35 40 45

Glu Thr Glu Glu Arg Arg Ile Asp Val Pro Leu Val Ile Lys Ala Lys  
50 55 60

Lys Asp Met Ala Asp Leu Gln Lys Leu Lys Glu Asp Leu Ala Asp Trp  
65 70 75 80

Leu Tyr Thr Glu Gln Pro Ala Glu Leu Ile Phe Asp Asp Glu Leu Asp  
85 90 95

Arg	Thr	Tyr	Leu	Ala	Leu	Ile	Asp	Gly	Ser	Val	Asp	Leu	Asp	Glu	Ile
			100					105					110		
Val	Asn	Arg	Gly	Arg	Gly	Val	Ile	Thr	Phe	Val	Cys	Pro	Met	Pro	Tyr
		115					120					125			
Lys	Leu	Gly	Lys	Thr	Asn	Thr	His	Lys	Phe	Thr	Gln	Glu	Trp	Ser	Thr
	130					135					140				
Glu	Thr	Thr	Ser	Tyr	Phe	Thr	Asn	Lys	Gly	Ser	Val	Glu	Ala	Pro	Ala
145					150					155					160
Leu	Ile	Glu	Met	Thr	Val	Lys	Lys	Pro	Ser	Thr	Phe	Leu	Asp	Val	Trp
				165					170					175	
Phe	Gly	Glu	Tyr	Pro	Asn	Asn	Arg	Asp	Tyr	Phe	Arg	Ile	Gly	Tyr	Pro
			180					185					190		
Leu	Thr	Val	Glu	Glu	Thr	Thr	Val	Gln	Glu	Arg	Glu	Arg	Val	Met	Trp
		195					200					205			
Asp	Glu	Met	Ala	Thr	Pro	Ile	Gly	Trp	Thr	Pro	Val	Thr	Gly	Gln	Phe
	210					215					220				
Asp	Asp	Met	Lys	Gly	Thr	Gly	Ser	Phe	Lys	Ser	Arg	Gly	Gly	Tyr	Ala
225					230					235					240
Leu	Tyr	Cys	Glu	Asp	Tyr	Gly	Lys	Asp	Val	Gly	Phe	Tyr	Gly	Ala	Ile
				245					250					255	
Ala	Lys	Lys	Asn	Ile	Pro	Gly	Gly	Pro	Leu	Gln	Asp	Phe	Glu	Met	Glu
			260					265					270		
Ala	Trp	Met	Thr	Leu	Lys	Ser	Lys	Asn	Ile	Gly	Glu	Met	Gly	Arg	Val
		275					280					285			
Glu	Val	Leu	Leu	Leu	Asp	Glu	Ala	Ser	Asn	Val	Val	Ala	Arg	Ile	Asn
	290					295					300				
Met	Asn	Asp	Leu	Tyr	Ala	Thr	Ala	Glu	Ile	Thr	Arg	Ala	His	Met	Lys
305					310					315					320
Ile	Gly	Asn	Ser	Gly	Thr	Pro	Asn	Ser	Phe	Arg	Lys	Leu	Val	Asp	Thr
				325					330					335	
Ser	Gly	Tyr	Tyr	Ser	Asn	Thr	Phe	Asn	Gln	Phe	Arg	Gly	Arg	Leu	Arg
			340					345					350		
Ile	Ala	Arg	Arg	Gly	Lys	Val	Trp	Ser	Val	Tyr	Val	Ala	Lys	Phe	Ile
		355					360					365			



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Asp Gly Thr Glu Lys Asp Gly Ala Ser Leu Val Glu Arg Trp Ile Asp  
370 375 380

Glu Thr Gly Asn Pro Met Thr Glu Arg Lys Ile Ala Gln Val Met Ile  
385 390 395 400

Ala Ile Cys Lys Trp Asp Asn His Gln Pro Val Asn Glu Ile Gln Ile  
405 410 415

Asp Asp Leu Lys Phe Trp Lys Val Asn Lys Val Pro Ser Asn Ala Gln  
420 425 430

Pro Tyr Ile Phe Asp Thr Gly Asp Lys Ile Val Ile Asp Thr Glu Lys  
435 440 445

Ser Leu Val Thr Ile Asn Gly Lys Asn Ala Ile Asn Ile Lys Glu Ile  
450 455 460

Phe Ser Asn Phe Pro Val Ile Ile Arg Gly Asp Asn Arg Ile Asp Ile  
465 470 475 480

Met Pro Pro Asp Val Asn Ala Thr Ile Ser Tyr Arg Glu Arg Tyr Arg  
485 490 495

<210> 30  
<211> 496  
<212> PRT  
<213> Bacillus anthracis

<400> 30

Met Gly Lys Leu Ser Phe Thr Phe Asn Asn Ile Arg Lys Asp Tyr Ile  
1 5 10 15

Gln Met Leu Val Gly Arg Lys Arg Pro Ser Trp Ala Pro Val Lys Arg  
20 25 30

Arg Leu Val Arg Val Pro His Arg Ala Gly Ala Leu Leu Leu Asn Thr  
35 40 45

Glu Thr Glu Glu Arg Arg Ile Asp Val Pro Leu Val Ile Lys Ala Lys  
50 55 60

Lys Asp Met Ala Asp Leu Gln Lys Leu Lys Glu Asp Leu Ala Asp Trp  
65 70 75 80

Leu Tyr Thr Glu Gln Pro Ala Glu Leu Ile Phe Asp Asp Glu Leu Asp  
85 90 95

Arg Thr Tyr Leu Ser Leu Ile Asp Gly Ser Val Asp Leu Asp Glu Ile  
100 105 110

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Val	Asn	Arg	Gly	Lys	Gly	Val	Ile	Thr	Phe	Val	Cys	Pro	Met	Pro	Tyr
		115					120					125			
Lys	Leu	Gly	Lys	Ile	Asn	Thr	His	Lys	Phe	Thr	Gln	Glu	Trp	Ser	Thr
	130					135					140				
Glu	Thr	Thr	Ser	Tyr	Phe	Thr	Asn	Lys	Gly	Ser	Val	Glu	Ala	Pro	Ala
145					150					155					160
Leu	Ile	Glu	Met	Thr	Val	Lys	Lys	Pro	Ser	Thr	Phe	Leu	Asp	Val	Trp
				165					170					175	
Phe	Gly	Glu	Tyr	Pro	His	Asn	Arg	Asp	Tyr	Phe	Arg	Ile	Gly	Tyr	Pro
			180					185					190		
Leu	Thr	Val	Glu	Glu	Thr	Thr	Val	Gln	Glu	Arg	Glu	Arg	Val	Met	Trp
		195					200					205			
Asp	Glu	Met	Ala	Thr	Pro	Ile	Gly	Trp	Thr	Pro	Val	Thr	Gly	Gln	Phe
	210					215					220				
Glu	Glu	Met	Lys	Gly	Thr	Gly	Ser	Phe	Lys	Ser	Arg	Gly	Gly	His	Ala
225					230					235					240
Leu	Tyr	Cys	Glu	Asp	Tyr	Gly	Lys	Glu	Thr	Gly	Phe	Tyr	Gly	Ala	Ile
				245					250					255	
Ala	Lys	Lys	Asn	Ile	Pro	Gly	Gly	Pro	Leu	Gln	Asp	Phe	Glu	Met	Glu
			260					265					270		
Ala	Trp	Val	Thr	Leu	Lys	Ser	Lys	Asn	Ile	Ser	Glu	Met	Gly	Arg	Val
		275					280					285			
Glu	Val	Leu	Leu	Leu	Asp	Glu	Thr	Ser	Asn	Val	Ile	Ser	Arg	Ile	Asn
	290					295					300				
Met	Asn	Asp	Leu	Tyr	Ala	Thr	Ala	Glu	Ile	Thr	Arg	Ala	His	Met	Thr
305					310					315					320
Ile	Gly	Asn	Ser	Gly	Thr	Pro	Asn	Ser	Phe	Arg	Lys	Leu	Val	Asp	Thr
				325					330					335	
Ser	Gly	Phe	Tyr	Ser	Thr	Thr	Phe	Asn	Gln	Phe	Arg	Gly	Arg	Leu	Arg
			340					345					350		
Ile	Ala	Arg	Arg	Gly	Lys	Val	Trp	Ser	Val	Tyr	Val	Ala	Lys	Phe	Ile
		355					360					365			
Asp	Gly	Thr	Glu	Lys	Asp	Gly	Ala	Ser	Leu	Val	Glu	Arg	Trp	Ile	Asp
	370					375					380				

Glu Thr Gly Asn Pro Met Thr Glu Arg Lys Ile Ala Gln Val Met Ile  
385 390 395 400

Ala Ile Cys Lys Trp Asp Asn His Gln Pro Ile Asn Glu Met Gln Ile  
405 410 415

Asp Asp Leu Lys Ile Trp Lys Val Asn Lys Val Pro Ser Asn Ala Gln  
420 425 430

Pro Tyr Ile Phe Asp Thr Gly Asp Lys Ile Val Ile Asp Thr Glu Lys  
435 440 445

Ser Leu Val Thr Ile Asn Gly Glu Lys Ala Ile Asn Ile Lys Glu Ile  
450 455 460

Phe Ser Asn Phe Pro Val Val Ile Arg Gly Glu Asn Arg Ile Asp Ile  
465 470 475 480

Met Pro Pro Asp Val Asn Ala Thr Ile Ser Tyr Arg Glu Arg Tyr Arg  
485 490 495

<210> 31  
<211> 1331  
<212> PRT  
<213> Bacillus anthracis

<400> 31

Met Arg Thr Pro Ser Gly Ile Leu His Val Val Asp Phe Lys Thr Asp  
1 5 10 15

Gln Ile Val Ala Ala Ile Gln Pro Glu Asp Tyr Trp Asp Asp Lys Arg  
20 25 30

His Trp Glu Leu Lys Asn Asn Val Asp Met Leu Asp Phe Thr Ala Phe  
35 40 45

Asp Gly Thr Asp His Ala Val Thr Leu Gln Gln Gln Asn Leu Val Leu  
50 55 60

Lys Glu Val Arg Asp Gly Arg Ile Val Pro Tyr Val Ile Thr Glu Thr  
65 70 75 80

Glu Lys Asn Ser Asp Thr Arg Ser Ile Thr Thr Tyr Ala Ser Gly Ala  
85 90 95

Trp Ile Gln Ile Ala Lys Ser Gly Ile Ile Lys Pro Gln Arg Ile Glu  
100 105 110

Ser Lys Thr Val Asn Glu Phe Met Asp Leu Ala Leu Leu Gly Met Lys  
115 120 125

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Trp Lys Arg Gly Ile Thr Glu Tyr Ala Gly Phe His Thr Met Thr Ile  
130 135 140

Asp Glu Tyr Ile Asp Pro Leu Thr Phe Leu Lys Lys Ile Ala Ser Leu  
145 150 155 160

Phe Lys Leu Glu Ile Arg Tyr Arg Val Glu Ile Lys Gly Ser Arg Ile  
165 170 175

Ile Gly Trp Tyr Val Asp Met Ile Gln Lys Arg Gly His Asp Thr Gly  
180 185 190

Lys Glu Ile Glu Leu Gly Lys Asp Leu Val Gly Val Thr Arg Ile Glu  
195 200 205

His Thr Arg Asn Ile Cys Ser Ala Leu Val Gly Phe Val Lys Gly Glu  
210 215 220

Gly Asp Lys Val Ile Thr Ile Glu Ser Ile Asn Lys Gly Leu Pro Tyr  
225 230 235 240

Ile Val Asp Ala Asp Ala Phe Gln Arg Trp Asn Glu His Gly Gln His  
245 250 255

Lys Phe Gly Phe Tyr Thr Pro Glu Thr Glu Glu Leu Asp Met Thr Pro  
260 265 270

Lys Arg Leu Leu Thr Leu Met Glu Ile Glu Leu Lys Lys Arg Val Asn  
275 280 285

Ser Ser Ile Ser Tyr Glu Val Glu Ala Gln Ser Ile Gly Arg Ile Phe  
290 295 300

Gly Leu Glu His Glu Leu Ile Asn Glu Gly Asp Thr Ile Lys Ile Lys  
305 310 315 320

Asp Thr Gly Phe Thr Pro Glu Leu Tyr Leu Glu Ala Arg Val Ile Ala  
325 330 335

Gly Asp Glu Ser Phe Thr Asp Ser Thr Gln Asp Lys Tyr Glu Phe Gly  
340 345 350

Asp Tyr Arg Glu Ile Val Asn Gln Asn Glu Glu Leu Arg Lys Ile Tyr  
355 360 365

Asn Arg Ile Leu Ser Ser Leu Gly Asn Lys Gln Glu Met Ile Asp Gln  
370 375 380

Leu Asp Arg Leu Val Gln Glu Ala Asn Glu Thr Ala Ser Asn Ala Lys

385

390

395

400

Lys Glu Ser Glu Ala Ala Lys Thr Leu Ala Glu Lys Val Gln Glu Asn  
405 410 415

Ile Lys Asn Asn Thr Val Glu Ile Ile Glu Ser Lys Asn Pro Pro Thr  
420 425 430

Thr Gly Leu Lys Pro Phe Lys Thr Leu Trp Arg Asp Ile Ser Ile Gly  
435 440 445

Lys Pro Gly Ile Leu Lys Ile Trp Thr Gly Thr Ala Trp Glu Ser Val  
450 455 460

Val Pro Asp Val Glu Ser Val Lys Lys Glu Thr Leu Asp Gln Val Asn  
465 470 475 480

Lys Asp Ile Ala Thr Thr Lys Thr Glu Leu Asn Gln Lys Val Gln Glu  
485 490 495

Ala Gln Asn Gln Ala Thr Gly Gln Phe Asn Glu Val Lys Glu Ser Leu  
500 505 510

Gln Gly Val Ser Arg Thr Ile Ser Asn Val Glu Asn Lys Gln Gly Glu  
515 520 525

Ile Asp Lys Lys Ile Thr Lys Phe Glu Gln Asp Ser Ser Gly Phe Lys  
530 535 540

Thr Ser Ile Glu Ser Leu Thr Lys Lys Asp Thr Glu Ile Ser Asn Lys  
545 550 555 560

Leu Asn Thr Val Glu Ser Thr Val Glu Gly Thr Lys Lys Thr Ile Ser  
565 570 575

Glu Val Gln Gln Thr Thr Asn Asp Leu Lys Lys Lys Thr Thr Glu Ile  
580 585 590

Glu Glu Lys Ala Gly Lys Ile Thr Glu Lys Leu Thr Ser Leu Glu Thr  
595 600 605

Arg Glu Val Asn Val Arg Asn Tyr Val Ile Asn Ser Asp Phe Ser Asn  
610 615 620

Val Thr Asn Ser Trp Ile Gly Ile Thr Asn Ala Thr Leu Phe Lys Phe  
625 630 635 640

Val Asp Val Asn Ile Ser Glu Ala Ser Ala Ile Lys Lys Gly Leu Gln  
645 650 655

PCT-US2005-009928\_Sequence Listing.txt as filed 10-3-05 in PCT  
Ile Thr Ser Asn Lys Ala Phe Val Tyr Gln Lys Leu Pro Ala Asp Val  
660 665 670

Phe Lys Lys Lys Lys Gly Ile Ala Ser Cys Tyr Ile Asn Val Ser Ser  
675 680 685

Phe Thr Pro Gly Thr Asp Tyr Pro Arg Leu Tyr Met Arg Phe Thr Tyr  
690 695 700

Asp Gln Asn Gly Thr Glu Lys Gln Tyr Tyr Ala Ile Leu Lys Gln Gln  
705 710 715 720

Glu Val Thr Asn Gly Trp Ile Arg Ile Ser Ile Pro Phe Asp Thr Thr  
725 730 735

Gly Tyr Thr Gly Glu Leu Lys Glu Val Arg Val Asn Ile Ala Thr Ala  
740 745 750

Asp Thr Thr Thr Ile Asp Ala Thr Phe Thr Gly Ile Met Val Thr Phe  
755 760 765

Gly Asp Leu Ile Glu Ser Trp Asn Leu Ala Pro Glu Asp Gly Val Thr  
770 775 780

Gln Gly Val Phe Gln Ser Lys Thr Thr Glu Ile Glu Lys Ser Val Asp  
785 790 795 800

Gly Val Lys Thr Thr Val Thr Asn Val Gln Asn Ser Gln Ala Gly Phe  
805 810 815

Glu Lys Arg Met Ser Asn Val Glu Gln Thr Ala Thr Gly Leu Ser Ser  
820 825 830

Thr Val Ser Asn Leu Asn Asn Val Val Ser Asp Gln Gly Lys Lys Leu  
835 840 845

Thr Glu Ala Asn Thr Lys Leu Glu Gln Gln Ala Thr Ala Ile Gly Ala  
850 855 860

Lys Val Glu Leu Lys Gln Val Glu Asp Tyr Val Ala Gly Phe Lys Ile  
865 870 875 880

Pro Glu Leu Lys Gln Thr Val Asp Lys Asn Lys Gln Asp Leu Leu Asp  
885 890 895

Glu Leu Ala Asn Lys Leu Ala Thr Glu Gln Phe Asn Gln Lys Met Thr  
900 905 910

Leu Ile Asp Asn Arg Phe Thr Ile Asn Glu Gln Gly Ile Asn Ala Ala  
915 920 925



Ala Lys Lys Thr Glu Val Tyr Thr Lys Thr Gln Ala Asp Gly Gln Phe  
930 935 940

Ala Thr Asp Ser Tyr Val Arg Asp Met Glu Ser Arg Leu Gln Leu Thr  
945 950 955 960

Glu Lys Gly Val Ser Ile Ser Val Lys Glu Asn Asp Val Ile Ala Ala  
965 970 975

Ile Asn Met Ser Lys Glu Asn Ile Lys Leu Asn Ala Ala Arg Ile Asp  
980 985 990

Leu Val Gly Lys Val Asn Ala Glu Trp Ile Lys Ala Gly Leu Leu Ser  
995 1000 1005

Gly Cys Gln Ile Arg Thr Ser Asn Thr Asp Asn Tyr Val Ser Leu  
1010 1015 1020

Asp Asp Gln Phe Ile Arg Leu Tyr Glu Arg Gly Val Ala Arg Ala  
1025 1030 1035

Phe Leu Gly His Tyr Arg Arg Ser Asp Gly Ala Val Gln Pro Thr  
1040 1045 1050

Phe Ile Leu Gly Ser Asp Glu Lys Thr Asn Ala Pro Glu Gly Thr  
1055 1060 1065

Leu Phe Met Ser Gln Ala Gly Ala Gly Trp Ser Gly Ala Tyr Ala  
1070 1075 1080

Ser Ile Gly Ile Ser Asn Gly Ile Val Asp Gly Ala Val Gln Lys  
1085 1090 1095

Ser Val Tyr Trp Glu Leu Gln Arg Asn Gly Leu Ser Val Leu Asn  
1100 1105 1110

Ala Asn Asp Tyr His Val Phe Tyr Ala Gly Asn Gly Asn Trp Tyr  
1115 1120 1125

Phe Arg Arg Gly Lys Pro Gly Leu Tyr Gln Thr Ser Leu Val Val  
1130 1135 1140

Glu Asp Asn Ser Thr Asp Ser Asp Leu Arg Leu Pro Asn Val Thr  
1145 1150 1155

Ile Arg Asn Ser Arg Ala Ala Gly Tyr Thr Gly Val Ile Gln Leu  
1160 1165 1170

Lys Ser Pro Val Thr Gln Asn Gly Trp Gly Ala Val Gln Gly Asn  
1175 1180 1185

PCT-US2005-009928\_Sequence Listing.txt as filed 10-3-05 in PCT

Phe Met Thr Pro Ser Leu Arg Glu Tyr Lys Ser Asn Ile Arg Asp  
1190 1195 1200

Ile Ser Phe Ser Ala Leu Glu Lys Ile Arg Ser Leu Lys Ile Arg  
1205 1210 1215

Gln Phe Asn Tyr Lys Asn Ala Val Asn Glu Leu Tyr Arg Met Arg  
1220 1225 1230

Glu Glu Lys Ser Pro Asn Asp Pro Pro Leu Thr Thr Glu Asp Ile  
1235 1240 1245

Lys Thr Tyr Tyr Gly Leu Ile Val Asp Glu Cys Asp Glu Met Phe  
1250 1255 1260

Val Asp Glu Ser Gly Lys Gly Ile His Leu Tyr Ser Tyr Ala Ser  
1265 1270 1275

Ile Gly Ile Lys Gly Leu Gln Glu Val Asp Ala Thr Val Gln Glu  
1280 1285 1290

Gln Glu Val Glu Ile Ala Asn Leu Lys Ser Gln Ile Ala Ser Gln  
1295 1300 1305

Glu Asp Arg Ile Ala Arg Leu Glu Glu Leu Leu Leu Gln Gln Leu  
1310 1315 1320

Ile Asn Lys Lys Pro Glu Gln Pro  
1325 1330

<210> 32  
<211> 1331  
<212> PRT  
<213> Bacillus anthracis

<400> 32

Met Arg Thr Pro Ser Gly Ile Leu His Val Val Asp Phe Lys Thr Asp  
1 5 10 15

Gln Ile Val Ala Ala Ile Gln Pro Glu Asp Tyr Trp Asp Asp Lys Arg  
20 25 30

His Trp Glu Leu Lys Asn Asn Val Asp Met Leu Asp Phe Thr Ala Phe  
35 40 45

Asp Gly Thr Asp His Ala Val Thr Leu Gln Gln Gln Asn Leu Val Leu  
50 55 60

Lys Glu Val Arg Asp Gly Arg Ile Val Pro Tyr Val Ile Thr Glu Thr  
65 70 75 80

PCT-US2005-009928\_Sequence Listing.txt as filed 10-3-05 in PCT

Glu	Lys	Asn	Ser	Asp	Thr	Arg	Ser	Ile	Thr	Thr	Tyr	Ala	Ser	Gly	Ala	
				85					90					95		
Trp	Ile	Gln	Ile	Ala	Lys	Ser	Gly	Ile	Ile	Lys	Pro	Gln	Arg	Ile	Glu	
			100					105					110			
Ser	Lys	Thr	Val	Asn	Glu	Phe	Met	Asp	Leu	Ala	Leu	Leu	Gly	Met	Lys	
		115					120					125				
Trp	Lys	Arg	Gly	Ile	Thr	Glu	Tyr	Ala	Gly	Phe	His	Thr	Met	Thr	Ile	
	130					135					140					
Asp	Glu	Tyr	Ile	Asp	Pro	Leu	Thr	Phe	Leu	Lys	Lys	Ile	Ala	Ser	Leu	
145					150					155					160	
Phe	Lys	Leu	Glu	Ile	Arg	Tyr	Arg	Val	Glu	Ile	Lys	Gly	Ser	Arg	Ile	
				165					170					175		
Ile	Gly	Trp	Tyr	Val	Asp	Met	Ile	Gln	Lys	Arg	Gly	His	Asp	Thr	Gly	
			180					185					190			
Lys	Glu	Ile	Glu	Leu	Gly	Lys	Asp	Leu	Val	Gly	Val	Thr	Arg	Ile	Glu	
		195					200					205				
His	Thr	Arg	Asn	Ile	Cys	Ser	Ala	Leu	Val	Gly	Phe	Val	Lys	Gly	Glu	
	210					215					220					
Gly	Asp	Lys	Val	Ile	Thr	Ile	Glu	Ser	Ile	Asn	Lys	Gly	Leu	Pro	Tyr	
225					230					235					240	
Ile	Val	Asp	Ala	Asp	Ala	Phe	Gln	Arg	Trp	Asn	Glu	His	Gly	Gln	His	
				245					250					255		
Lys	Phe	Gly	Phe	Tyr	Thr	Pro	Glu	Thr	Glu	Glu	Leu	Asp	Met	Thr	Pro	
			260					265					270			
Lys	Arg	Leu	Leu	Thr	Leu	Met	Glu	Ile	Glu	Leu	Lys	Lys	Arg	Val	Asn	
		275					280					285				
Ser	Ser	Ile	Ser	Tyr	Glu	Val	Glu	Ala	Gln	Ser	Ile	Gly	Arg	Ile	Phe	
	290					295					300					
Gly	Leu	Glu	His	Glu	Leu	Ile	Asn	Glu	Gly	Asp	Thr	Ile	Lys	Ile	Lys	
305					310					315					320	
Asp	Thr	Gly	Phe	Thr	Pro	Glu	Leu	Tyr	Leu	Glu	Ala	Arg	Val	Ile	Ala	
				325					330					335		
Gly	Asp	Glu	Ser	Phe	Thr	Asp	Ser	Thr	Gln	Asp	Lys	Tyr	Glu	Phe	Gly	

Asp Tyr Arg Glu Ile Val Asn Gln Asn Glu Glu Leu Arg Lys Ile Tyr  
355 360 365

Asn Arg Ile Leu Ser Ser Leu Gly Asn Lys Gln Glu Met Ile Asp Gln  
370 375 380

Leu Asp Arg Leu Val Gln Glu Ala Asn Glu Thr Ala Ser Asn Ala Lys  
385 390 395 400

Lys Glu Ser Glu Ala Ala Lys Thr Leu Ala Glu Lys Val Gln Glu Asn  
405 410 415

Ile Lys Asn Asn Thr Val Glu Ile Ile Glu Ser Lys Asn Pro Pro Thr  
420 425 430

Thr Gly Leu Lys Pro Phe Lys Thr Leu Trp Arg Asp Ile Ser Ile Gly  
435 440 445

Lys Pro Gly Ile Leu Lys Ile Trp Thr Gly Thr Ala Trp Glu Ser Val  
450 455 460

Val Pro Asp Val Glu Ser Val Lys Lys Glu Thr Leu Asp Gln Val Asn  
465 470 475 480

Lys Asp Ile Ala Thr Thr Lys Thr Glu Leu Asn Gln Lys Val Gln Glu  
485 490 495

Ala Gln Asn Gln Ala Thr Gly Gln Phe Asn Glu Val Lys Glu Ser Leu  
500 505 510

Gln Gly Val Ser Arg Thr Ile Ser Asn Val Glu Asn Lys Gln Gly Glu  
515 520 525

Ile Asp Lys Lys Ile Thr Lys Phe Glu Gln Asp Ser Ser Gly Phe Lys  
530 535 540

Thr Ser Ile Glu Ser Leu Thr Lys Lys Asp Thr Glu Ile Ser Asn Lys  
545 550 555 560

Leu Asn Thr Val Glu Ser Thr Val Glu Gly Thr Lys Lys Thr Ile Ser  
565 570 575

Glu Val Gln Gln Thr Thr Asn Asp Leu Lys Lys Lys Thr Thr Glu Ile  
580 585 590

Glu Glu Lys Ala Gly Lys Ile Thr Glu Lys Leu Thr Ser Leu Glu Thr  
595 600 605

PCT-US2005-009928\_Sequence Listing.txt as filed 10-3-05 in PCT  
 Arg Glu Val Asn Val Arg Asn Tyr Val Ile Asn Ser Asp Phe Ser Asn  
 610 615 620

Val Thr Asn Ser Trp Ile Gly Ile Thr Asn Ala Thr Leu Phe Lys Phe  
 625 630 635 640

Val Asp Val Asn Ile Ser Glu Ala Ser Ala Ile Lys Lys Gly Leu Gln  
 645 650 655

Ile Thr Ser Asn Lys Ala Phe Val Tyr Gln Lys Leu Pro Ala Asp Val  
 660 665 670

Phe Lys Lys Lys Lys Gly Ile Ala Ser Cys Tyr Ile Asn Val Ser Ser  
 675 680 685

Phe Thr Pro Gly Thr Asp Tyr Pro Arg Leu Tyr Met Arg Phe Thr Tyr  
 690 695 700

Asp Gln Asn Gly Thr Glu Lys Gln Tyr Tyr Ala Ile Leu Lys Gln Gln  
 705 710 715 720

Glu Val Thr Asn Gly Trp Ile Arg Ile Ser Ile Pro Phe Asp Thr Thr  
 725 730 735

Gly Tyr Thr Gly Glu Leu Lys Glu Val Arg Val Asn Ile Ala Thr Ala  
 740 745 750

Asp Thr Thr Thr Ile Asp Ala Thr Phe Thr Gly Ile Met Val Thr Phe  
 755 760 765

Gly Asp Leu Ile Glu Ser Trp Asn Leu Ala Pro Glu Asp Gly Val Thr  
 770 775 780

Gln Gly Val Phe Gln Ser Lys Thr Thr Glu Ile Glu Lys Ser Val Asp  
 785 790 795 800

Gly Val Lys Thr Thr Val Thr Asn Val Gln Asn Ser Gln Ala Gly Phe  
 805 810 815

Glu Lys Arg Met Ser Asn Val Glu Gln Thr Ala Thr Gly Leu Ser Ser  
 820 825 830

Thr Val Ser Asn Leu Asn Asn Val Val Ser Asp Gln Gly Lys Lys Leu  
 835 840 845

Thr Glu Ala Asn Thr Lys Leu Glu Gln Gln Ala Thr Ala Ile Gly Ala  
 850 855 860

Lys Val Glu Leu Lys Gln Val Glu Asp Tyr Val Ala Gly Phe Lys Ile  
 865 870 875 880

PCT-US2005-009928\_Sequence Listing.txt as filed 10-3-05 in PCT

Pro Glu Leu Lys Gln Thr Val Asp Lys Asn Lys Gln Asp Leu Leu Asp  
885 890 895

Glu Leu Ala Asn Lys Leu Ala Thr Glu Gln Phe Asn Gln Lys Met Thr  
900 905 910

Leu Ile Asp Asn Arg Phe Thr Ile Asn Glu Gln Gly Ile Asn Ala Ala  
915 920 925

Ala Lys Lys Thr Glu Val Tyr Thr Lys Thr Gln Ala Asp Gly Gln Phe  
930 935 940

Ala Thr Asp Ser Tyr Val Arg Asp Met Glu Ser Arg Leu Gln Leu Thr  
945 950 955 960

Glu Lys Gly Val Ser Ile Ser Val Lys Glu Asn Asp Val Ile Ala Ala  
965 970 975

Ile Asn Met Ser Lys Glu Asn Ile Lys Leu Asn Ala Ala Arg Ile Asp  
980 985 990

Leu Val Gly Lys Val Asn Ala Glu Trp Ile Lys Ala Gly Leu Leu Ser  
995 1000 1005

Gly Cys Gln Ile Arg Thr Ser Asn Thr Asp Asn Tyr Val Ser Leu  
1010 1015 1020

Asp Asp Gln Phe Ile Arg Leu Tyr Glu Arg Gly Val Ala Arg Ala  
1025 1030 1035

Phe Leu Gly His Tyr Arg Arg Ser Asp Gly Ala Val Gln Pro Thr  
1040 1045 1050

Phe Ile Leu Gly Ser Asp Glu Lys Thr Asn Ala Pro Glu Gly Thr  
1055 1060 1065

Leu Phe Met Ser Gln Ala Gly Ala Gly Trp Ser Gly Ala Tyr Ala  
1070 1075 1080

Ser Ile Gly Ile Ser Asn Gly Ile Val Asp Gly Ala Val Gln Lys  
1085 1090 1095

Ser Val Tyr Trp Glu Leu Gln Arg Asn Gly Leu Ser Val Leu Asn  
1100 1105 1110

Ala Asn Asp Tyr His Val Phe Tyr Ala Gly Asn Gly Asn Trp Tyr  
1115 1120 1125

Phe Arg Arg Gly Lys Pro Gly Leu Tyr Gln Thr Ser Leu Val Val  
1130 1135 1140



PCT-US2005-009928\_Sequence Listing.txt as filed 10-3-05 in PCT

Glu Asp Asn Ser Thr Asp Ser Asp Leu Arg Leu Pro Asn Val Thr  
1145 1150 1155

Ile Arg Asn Ser Arg Ala Ala Gly Tyr Thr Gly Val Ile Gln Leu  
1160 1165 1170

Lys Ser Pro Val Thr Gln Asn Gly Trp Gly Ala Val Gln Gly Asn  
1175 1180 1185

Phe Met Thr Pro Ser Leu Arg Glu Tyr Lys Ser Asn Ile Arg Asp  
1190 1195 1200

Ile Ser Phe Ser Ala Leu Glu Lys Ile Arg Ser Leu Lys Ile Arg  
1205 1210 1215

Gln Phe Asn Tyr Lys Asn Ala Val Asn Glu Leu Tyr Arg Met Arg  
1220 1225 1230

Glu Glu Lys Ser Pro Asn Asp Pro Pro Leu Thr Thr Glu Asp Ile  
1235 1240 1245

Lys Thr Tyr Tyr Gly Leu Ile Val Asp Glu Cys Asp Glu Met Phe  
1250 1255 1260

Val Asp Glu Ser Gly Lys Gly Ile His Leu Tyr Ser Tyr Ala Ser  
1265 1270 1275

Ile Gly Ile Lys Gly Leu Gln Glu Val Asp Ala Thr Val Gln Glu  
1280 1285 1290

Gln Glu Val Glu Ile Ala Asn Leu Lys Ser Gln Ile Ala Ser Gln  
1295 1300 1305

Glu Asp Arg Ile Ala Arg Leu Glu Glu Leu Leu Leu Gln Gln Leu  
1310 1315 1320

Ile Asn Lys Lys Pro Glu Gln Pro  
1325 1330

<210> 33  
<211> 141  
<212> PRT  
<213> Bacillus anthracis

<400> 33

Met Asp Arg Ile Asp Val Leu Leu Lys Ala Phe Ile Ala Ala Phe Gly  
1 5 10 15

Gly Phe Cys Gly Tyr Phe Leu Gly Gly Trp Asp Ala Thr Leu Lys Ile  
20 25 30

PCT-US2005-009928\_Sequence Listing.txt as filed 10-3-05 in PCT

Leu Val Thr Met Val Val Ile Asp Tyr Leu Thr Gly Met Ile Ala Ala  
35 40 45

Gly Tyr Asn Gly Glu Leu Lys Ser Lys Val Gly Phe Lys Gly Ile Ala  
50 55 60

Lys Lys Val Val Leu Phe Leu Leu Val Gly Ala Ala Ala Gln Leu Asp  
65 70 75 80

Ser Ala Leu Gly Ser Asn Ser Ala Ile Arg Glu Ala Thr Ile Phe Phe  
85 90 95

Phe Met Gly Asn Glu Leu Leu Ser Leu Leu Glu Asn Ala Gly Arg Met  
100 105 110

Gly Ile Pro Leu Pro Gln Ala Leu Thr Asn Ala Val Glu Ile Leu Gly  
115 120 125

Gly Lys Gln Lys Gln Glu Glu Lys Lys Gly Asp Val Gln  
130 135 140

<210> 34  
<211> 141  
<212> PRT  
<213> Bacillus anthracis

<400> 34

Met Asp Arg Ile Asp Val Leu Leu Lys Ala Phe Ile Ala Ala Phe Gly  
1 5 10 15

Gly Phe Cys Gly Tyr Phe Leu Gly Gly Trp Asp Ala Thr Leu Lys Ile  
20 25 30

Leu Val Thr Met Val Val Ile Asp Tyr Leu Thr Gly Met Ile Ala Ala  
35 40 45

Gly Tyr Asn Gly Glu Leu Lys Ser Lys Val Gly Phe Lys Gly Ile Ala  
50 55 60

Lys Lys Val Val Leu Phe Leu Leu Val Gly Ala Ala Ala Gln Leu Asp  
65 70 75 80

Ser Ala Leu Gly Ser Asn Ser Ala Ile Arg Glu Ala Thr Ile Phe Phe  
85 90 95

Phe Met Gly Asn Glu Leu Leu Ser Leu Leu Glu Asn Ala Gly Arg Met  
100 105 110

Gly Ile Pro Leu Pro Gln Ala Leu Thr Asn Ala Val Glu Ile Leu Gly  
115 120 125

PCT-US2005-009928\_Sequence Listing.txt as filed 10-3-05 in PCT

Gly Lys Gln Lys Gln Glu Glu Lys Lys Gly Asp Val Gln  
130 135 140

<210> 35  
<211> 233  
<212> PRT  
<213> Bacillus anthracis  
<400> 35

Met Glu Ile Gln Lys Lys Leu Val Asp Pro Ser Lys Tyr Gly Thr Lys  
1 5 10 15

Cys Pro Tyr Thr Met Lys Pro Lys Tyr Ile Thr Val His Asn Thr Tyr  
20 25 30

Asn Asp Ala Pro Ala Glu Asn Glu Val Ser Tyr Met Ile Ser Asn Asn  
35 40 45

Asn Glu Val Ser Phe His Ile Ala Val Asp Asp Lys Lys Ala Ile Gln  
50 55 60

Gly Ile Pro Leu Glu Arg Asn Ala Trp Ala Cys Gly Asp Gly Asn Gly  
65 70 75 80

Ser Gly Asn Arg Gln Ser Ile Ser Val Glu Ile Cys Tyr Ser Lys Ser  
85 90 95

Gly Gly Asp Arg Tyr Tyr Lys Ala Glu Asp Asn Ala Val Asp Val Val  
100 105 110

Arg Gln Leu Met Ser Met Tyr Asn Ile Pro Ile Glu Asn Val Arg Thr  
115 120 125

His Gln Ser Trp Ser Gly Lys Tyr Cys Pro His Arg Met Leu Ala Glu  
130 135 140

Gly Arg Trp Gly Ala Phe Ile Gln Lys Val Lys Asn Gly Asn Val Ala  
145 150 155 160

Thr Thr Ser Pro Thr Lys Gln Asn Ile Ile Gln Ser Gly Ala Phe Ser  
165 170 175

Pro Tyr Glu Thr Pro Asp Val Met Gly Ala Leu Thr Ser Leu Lys Met  
180 185 190

Thr Ala Asp Phe Ile Leu Gln Ser Asp Gly Leu Thr Tyr Phe Ile Ser  
195 200 205

Lys Pro Thr Ser Asp Ala Gln Leu Lys Ala Met Lys Glu Tyr Leu Asp  
210 215 220

Arg Lys Gly Trp Trp Tyr Glu Val Lys  
225 230

<210> 36  
<211> 233  
<212> PRT  
<213> Bacillus anthracis  
<400> 36

Met Glu Ile Gln Lys Lys Leu Val Asp Pro Ser Lys Tyr Gly Thr Lys  
1 5 10 15

Cys Pro Tyr Thr Met Lys Pro Lys Tyr Ile Thr Val His Asn Thr Tyr  
20 25 30

Asn Asp Ala Pro Ala Glu Asn Glu Val Ser Tyr Met Ile Ser Asn Asn  
35 40 45

Asn Glu Val Ser Phe His Ile Ala Val Asp Asp Lys Lys Ala Ile Gln  
50 55 60

Gly Ile Pro Leu Glu Arg Asn Ala Trp Ala Cys Gly Asp Gly Asn Gly  
65 70 75 80

Ser Gly Asn Arg Gln Ser Ile Ser Val Glu Ile Cys Tyr Ser Lys Ser  
85 90 95

Gly Gly Asp Arg Tyr Tyr Lys Ala Glu Asp Asn Ala Val Asp Val Val  
100 105 110

Arg Gln Leu Met Ser Met Tyr Asn Ile Pro Ile Glu Asn Val Arg Thr  
115 120 125

His Gln Ser Trp Ser Gly Lys Tyr Cys Pro His Arg Met Leu Ala Glu  
130 135 140

Gly Arg Trp Gly Ala Phe Ile Gln Lys Val Lys Asn Gly Asn Val Ala  
145 150 155 160

Thr Thr Ser Pro Thr Lys Gln Asn Ile Ile Gln Ser Gly Ala Phe Ser  
165 170 175

Pro Tyr Glu Thr Pro Asp Val Met Gly Ala Leu Thr Ser Leu Lys Met  
180 185 190

Thr Ala Asp Phe Ile Leu Gln Ser Asp Gly Leu Thr Tyr Phe Ile Ser  
195 200 205

Lys Pro Thr Ser Asp Ala Gln Leu Lys Ala Met Lys Glu Tyr Leu Asp  
210 215 220

Arg Lys Gly Trp Trp Tyr Glu Val Lys  
225 230

<210> 37  
<211> 165  
<212> PRT  
<213> Bacillus anthracis

<400> 37

Met Lys Met Tyr Lys Lys Leu Ile Ser Ile Cys Ile Gly Ser Thr Leu  
1 5 10 15

Leu Leu Gly Leu Thr Ala Cys Asp Ser Ser Lys Gln Ser Glu Ser Ser  
20 25 30

Glu Lys Thr Asn Val Lys Ser Gln Pro Glu Thr Lys Lys Asp Leu Thr  
35 40 45

Ser Gln Asp Glu Leu Asn Lys Lys Ile Lys Gln Asp Ala Glu Glu Val  
50 55 60

Ser Phe Val Lys Ala Asn Gly Asp Gln Tyr Glu Lys Gly Lys Arg Leu  
65 70 75 80

Lys Ala Thr Gly Thr Val Asp Leu Leu Leu Lys Ser Ser Ala Leu Pro  
85 90 95

Ser Phe Val Ile Ser Thr Asn Glu Asn Asp Gly Lys Gly Met Tyr Thr  
100 105 110

Ile Gln Ile Val Gln Ser Gly Val Gln Thr Asn Glu Asn Glu Ile Thr  
115 120 125

Leu Lys Asn Gly Leu Lys Ile Ser Lys Gly Ser Ile Val Thr Ile Tyr  
130 135 140

Gly Ala Tyr Asp Glu Lys Asp Lys Thr Gly Met Pro Lys Ile Ser Ala  
145 150 155 160

Thr val Ile Glu Gln  
165

<210> 38  
<211> 165  
<212> PRT  
<213> Bacillus anthracis

<400> 38

Met Lys Met Tyr Lys Lys Leu Ile Ser Ile Cys Ile Gly Ser Thr Leu  
1 5 10 15

Leu Leu Gly Leu Thr Ala Cys Asp Ser Ser Lys Gln Ser Glu Ser Ser  
20 25 30

Glu Lys Thr Asn Val Lys Ser Gln Pro Glu Thr Lys Lys Asp Leu Thr  
35 40 45

Ser Gln Asp Glu Leu Asn Lys Lys Ile Lys Gln Asp Ala Glu Glu Val  
50 55 60

Ser Phe Val Lys Ala Asn Gly Asp Gln Tyr Glu Lys Gly Lys Arg Leu  
65 70 75 80

Lys Ala Thr Gly Thr Val Asp Leu Leu Leu Lys Ser Ser Ala Leu Pro  
85 90 95

Ser Phe Val Ile Ser Thr Asn Glu Asn Asp Gly Lys Gly Met Tyr Thr  
100 105 110

Ile Gln Ile Val Gln Ser Gly Val Gln Thr Asn Glu Asn Glu Ile Thr  
115 120 125

Leu Lys Asn Gly Leu Lys Ile Ser Lys Gly Ser Ile Val Thr Ile Tyr  
130 135 140

Gly Ala Tyr Asp Glu Lys Asp Lys Thr Gly Met Pro Lys Ile Ser Ala  
145 150 155 160

Thr Val Ile Glu Gln  
165

<210> 39  
<211> 70  
<212> PRT  
<213> Bacillus anthracis

<400> 39

Val Arg Leu Lys Cys Lys Leu Arg Val Ile Phe Ala Glu Arg Glu Ile  
1 5 10 15

Arg Gln Lys Glu Phe Ser Lys Leu Ile Gly Ile Ser Gln Thr Thr Met  
20 25 30

Ser Ser Leu Val Asn Asn Thr Thr Leu Pro Thr Phe Leu Thr Ala Tyr  
35 40 45

Lys Ile Ala Lys Glu Leu Lys Leu His Met Glu Glu Ile Trp Ile Glu  
50 55 60

Glu Glu Asn Glu Asn Val  
65 70



PCT-US2005-009928\_Sequence Listing.txt as filed 10-3-05 in PCT

<210> 40  
 <211> 70  
 <212> PRT  
 <213> Bacillus anthracis

<400> 40

Val Arg Leu Lys Cys Lys Leu Arg Val Ile Phe Ala Glu Arg Glu Ile  
 1 5 10 15

Arg Gln Lys Glu Phe Ser Lys Leu Ile Gly Ile Ser Gln Thr Thr Met  
 20 25 30

Ser Ser Leu Val Asn Asn Thr Thr Leu Pro Thr Phe Leu Thr Ala Tyr  
 35 40 45

Lys Ile Ala Lys Glu Leu Lys Leu His Met Glu Glu Ile Trp Ile Glu  
 50 55 60

Glu Glu Asn Glu Asn Val  
 65 70

<210> 41  
 <211> 102  
 <212> PRT  
 <213> Bacillus anthracis

<400> 41

Met Arg Trp Gln Tyr Asn His Leu Asn Thr Thr Pro Tyr Leu His Pro  
 1 5 10 15

Ser Lys Glu Leu Cys Ser Met Tyr Asn Gly Ser Arg Ser Arg Ala Glu  
 20 25 30

Thr Glu Ser Ile Leu Asn His Met Lys Asn His Glu Val Tyr Asp Arg  
 35 40 45

Lys Glu Tyr Lys Gly Tyr Phe Ser Leu Ser Gln Val Leu Glu Glu Asp  
 50 55 60

Leu Tyr Gly Glu Glu Glu Asp Val Leu Asn Trp Glu Ile Leu Met Asp  
 65 70 75 80

Cys Tyr Asp Val Val Leu Thr Arg Lys Gly Ile Ala Phe Arg Glu Lys  
 85 90 95

Glu Glu Glu Glu Gln Ala  
 100

<210> 42  
 <211> 102  
 <212> PRT  
 <213> Bacillus anthracis

<400> 42

Met Arg Trp Gln Tyr Asn His Leu Asn Thr Thr Pro Tyr Leu His Pro  
1 5 10 15  
Ser Lys Glu Leu Cys Ser Met Tyr Asn Gly Ser Arg Ser Arg Ala Glu  
20 25 30  
Thr Glu Ser Ile Leu Asn His Met Lys Asn His Glu Val Tyr Asp Arg  
35 40 45  
Lys Glu Tyr Lys Gly Tyr Phe Ser Leu Ser Gln Val Leu Glu Glu Asp  
50 55 60  
Leu Tyr Gly Glu Glu Glu Asp Val Leu Asn Trp Glu Ile Leu Met Asp  
65 70 75 80  
Cys Tyr Asp Val Val Leu Thr Arg Lys Gly Ile Ala Phe Arg Glu Lys  
85 90 95  
Glu Glu Glu Glu Gln Ala  
100

<210> 43

<211> 60

<212> PRT

<213> Bacillus anthracis

<400> 43

Met Thr Leu Ala Gly Glu Ala Ile Ile Ile Trp Thr Ala Thr Gly Leu  
1 5 10 15  
Ser Val Val Ala Met Lys Ala Ala Glu Lys Met Gly Lys Ser Val Pro  
20 25 30  
His Trp Leu Pro Arg Val Thr Leu Tyr Thr Thr Leu Thr Gly Ser Phe  
35 40 45  
Leu Tyr Leu Leu Arg Tyr Val Leu Val Leu Phe Leu  
50 55 60

<210> 44

<211> 60

<212> PRT

<213> Bacillus anthracis

<400> 44

Met Thr Leu Ala Gly Glu Ala Ile Ile Ile Trp Thr Ala Thr Gly Leu  
1 5 10 15  
Ser Val Val Ala Met Lys Ala Ala Glu Lys Met Gly Lys Ser Val Pro  
20 25 30

His Trp Leu Pro Arg Val Thr Leu Tyr Thr Thr Leu Thr Gly Ser Phe  
35 40 45

Leu Tyr Leu Leu Arg Tyr Val Leu Val Leu Phe Leu  
50 55 60

<210> 45  
<211> 429  
<212> PRT  
<213> Bacillus anthracis

<400> 45

Met Trp Lys Leu Phe Ile Pro Tyr Val Ile Arg Ser Leu Ala Cys Met  
1 5 10 15

His Val Phe Leu Glu Thr Gly Ile Tyr Thr Leu Tyr Lys Arg Asp Ile  
20 25 30

Arg Ser Asp Phe Met Leu Glu Leu Leu Ser Val Pro Phe Ala Gly Leu  
35 40 45

Ile Phe Ala Ile Val Gly Glu Arg Leu Lys Gly Arg Glu Ser Asp Arg  
50 55 60

Lys Lys Ile Gln Val Phe Phe Glu Val Ser Gly Ile Ala Ile Arg Arg  
65 70 75 80

Glu Asp Lys Leu Gln Tyr Pro Val Phe Leu Glu Gln Lys Glu Asp Asp  
85 90 95

Arg Ser Thr Thr Tyr Ile Tyr Arg Leu Pro Val Gly Met Pro Ser Lys  
100 105 110

Ile Ile Gln Lys Val Glu Asp Val Val Ser Glu Gly Leu Ser Lys Pro  
115 120 125

Val Arg Ile Asp Tyr Asp Asn Tyr Lys Leu Asn Ile Arg Val Phe His  
130 135 140

Arg Asp Ile Pro Lys Lys Trp Ser Trp Ser Lys Gly Leu Val Ala Glu  
145 150 155 160

Gly Ser Trp Cys Val Pro Met Gly Gln Ser Leu Glu Lys Leu Ile Tyr  
165 170 175

His Asp Phe Asp Lys Thr Pro His Met Thr Leu Gly Gly Leu Thr Arg  
180 185 190

Met Gly Lys Thr Val Phe Leu Lys Asn Val Val Thr Ser Leu Thr Leu  
195 200 205

PCT-US2005-009928\_Sequence Listing.txt as filed 10-3-05 in PCT

Ala Gln Pro Glu His Ile Asn Leu Tyr Ile Ile Asp Leu Lys Gly Gly  
210 215 220

Leu Glu Phe Gly Pro Tyr Lys Asn Leu Lys Gln Val Val Ser Ile Ala  
225 230 235 240

Glu Lys Pro Ala Glu Ala Phe Met Ile Leu Thr Asn Ile Leu Lys Lys  
245 250 255

Met Glu Glu Lys Met Glu Tyr Met Lys Cys Arg His Tyr Thr Asn Val  
260 265 270

Val Glu Thr Asn Ile Lys Glu Arg Tyr Phe Ile Ile Val Asp Glu Gly  
275 280 285

Ala Glu Leu Cys Pro Asp Lys Ser Met Lys Lys Glu Gln Gln Arg Leu  
290 295 300

Leu Gly Ala Cys Gln Gln Met Leu Ser His Ile Ala Arg Ile Gly Gly  
305 310 315 320

Ala Leu Gly Phe Arg Leu Ile Phe Cys Thr Gln Tyr Pro Thr Gly Asp  
325 330 335

Thr Leu Pro Arg Gln Val Lys Gln Asn Ser Asp Ala Lys Leu Gly Phe  
340 345 350

Arg Leu Pro Thr Gln Thr Ala Ser Ser Val Val Ile Asp Glu Ala Gly  
355 360 365

Leu Glu Thr Ile Lys Ser Ile Pro Gly Arg Ala Ile Phe Lys Thr Asp  
370 375 380

Arg Leu Thr Glu Ile Gln Val Pro Tyr Ile Ser Asn Glu Met Met Trp  
385 390 395 400

Glu His Leu Lys Gly Tyr Glu Val Glu Lys His Glu Asp Ala Asn Ala  
405 410 415

Tyr Ala Asn Gln Pro Ser Asn Gly Asp Thr Cys Asp Asp  
420 425

<210> 46  
<211> 429  
<212> PRT  
<213> Bacillus anthracis

<400> 46

Met Trp Lys Leu Phe Ile Pro Tyr Val Ile Arg Ser Leu Ala Cys Met  
1 5 10 15

PCT-US2005-009928\_Sequence Listing.txt as filed 10-3-05 in PCT

His Val Phe Leu Glu Thr Gly Ile Tyr Thr Leu Tyr Lys Arg Asp Ile  
20 25 30

Arg Ser Asp Phe Met Leu Glu Leu Leu Ser Val Pro Phe Ala Gly Leu  
35 40 45

Ile Phe Ala Ile Val Gly Glu Arg Leu Lys Gly Arg Glu Ser Asp Arg  
50 55 60

Lys Lys Ile Gln Val Phe Phe Glu Val Ser Gly Ile Ala Ile Arg Arg  
65 70 75 80

Glu Asp Lys Leu Gln Tyr Pro Val Phe Leu Glu Gln Lys Glu Asp Asp  
85 90 95

Arg Ser Thr Thr Tyr Ile Tyr Arg Leu Pro Val Gly Met Pro Ser Lys  
100 105 110

Ile Ile Gln Lys Val Glu Asp Val Val Ser Glu Gly Leu Ser Lys Pro  
115 120 125

Val Arg Ile Asp Tyr Asp Asn Tyr Lys Leu Asn Ile Arg Val Phe His  
130 135 140

Arg Asp Ile Pro Lys Lys Trp Ser Trp Ser Lys Gly Leu Val Ala Glu  
145 150 155 160

Gly Ser Trp Cys Val Pro Met Gly Gln Ser Leu Glu Lys Leu Ile Tyr  
165 170 175

His Asp Phe Asp Lys Thr Pro His Met Thr Leu Gly Gly Leu Thr Arg  
180 185 190

Met Gly Lys Thr Val Phe Leu Lys Asn Val Val Thr Ser Leu Thr Leu  
195 200 205

Ala Gln Pro Glu His Ile Asn Leu Tyr Ile Ile Asp Leu Lys Gly Gly  
210 215 220

Leu Glu Phe Gly Pro Tyr Lys Asn Leu Lys Gln Val Val Ser Ile Ala  
225 230 235 240

Glu Lys Pro Ala Glu Ala Phe Met Ile Leu Thr Asn Ile Leu Lys Lys  
245 250 255

Met Glu Glu Lys Met Glu Tyr Met Lys Cys Arg His Tyr Thr Asn Val  
260 265 270

Val Glu Thr Asn Ile Lys Glu Arg Tyr Phe Ile Ile Val Asp Glu Gly  
Page 96

Ala Glu Leu Cys Pro Asp Lys Ser Met Lys Lys Glu Gln Gln Arg Leu  
290 295 300

Leu Gly Ala Cys Gln Gln Met Leu Ser His Ile Ala Arg Ile Gly Gly  
305 310 315 320

Ala Leu Gly Phe Arg Leu Ile Phe Cys Thr Gln Tyr Pro Thr Gly Asp  
325 330 335

Thr Leu Pro Arg Gln Val Lys Gln Asn Ser Asp Ala Lys Leu Gly Phe  
340 345 350

Arg Leu Pro Thr Gln Thr Ala Ser Ser Val Val Ile Asp Glu Ala Gly  
355 360 365

Leu Glu Thr Ile Lys Ser Ile Pro Gly Arg Ala Ile Phe Lys Thr Asp  
370 375 380

Arg Leu Thr Glu Ile Gln Val Pro Tyr Ile Ser Asn Glu Met Met Trp  
385 390 395 400

Glu His Leu Lys Gly Tyr Glu Val Glu Lys His Glu Asp Ala Asn Ala  
405 410 415

Tyr Ala Asn Gln Pro Ser Asn Gly Asp Thr Cys Asp Asp  
420 425

<210> 47  
<211> 210  
<212> PRT  
<213> Bacillus anthracis

<400> 47

Met Arg Trp Arg Asn Met Arg Met Gln Thr His Met Gln Ile Asn Arg  
1 5 10 15

Gln Met Ala Ile Leu Ala Thr Ile Arg Lys Leu Gln Phe Ala Thr Arg  
20 25 30

Arg His Leu Met Ser Ile His Glu Met Gly Gly Ile Arg Asn Ala Asn  
35 40 45

Arg Ile Leu Lys Asp Leu Ser Ile Tyr Thr Ser Lys Val Val Tyr Asn  
50 55 60

Lys Glu His Val Tyr Tyr Leu Asn Gln Ser Gly His Lys Leu Phe Gly  
65 70 75 80

Glu Gly Lys Val Val His His Gly Lys Val Thr His Ala Leu Leu Arg



Asn Glu Ala Trp Leu Asn Leu Tyr Cys Pro Asp Asp Trp Gln Val Glu  
100 105 110

Thr Glu Ile Lys Tyr Ile Lys Asp Asn Lys Lys Lys Lys Ile Ile Pro  
115 120 125

Asp Val Lys Phe Arg Asp Glu Asp Arg Ile Leu His Ala Val Glu Ile  
130 135 140

Asp Arg Thr Gln Lys Met Ile Val Asn Asp Glu Lys Leu Lys Lys Tyr  
145 150 155 160

Glu Glu Leu Thr Gln Ile Tyr Lys Gln Lys His Asn Gly Lys Val Pro  
165 170 175

Val Ile His Phe Phe Thr Ile Thr Lys Tyr Arg Glu Lys Lys Leu Glu  
180 185 190

Glu Leu Ala Asn Lys Tyr Asn Val Phe Val Lys Val Tyr Val Ile Ala  
195 200 205

Thr Thr  
210

<210> 48  
<211> 210  
<212> PRT  
<213> Bacillus anthracis

<400> 48

Met Arg Trp Arg Asn Met Arg Met Gln Thr His Met Gln Ile Asn Arg  
1 5 10 15

Gln Met Ala Ile Leu Ala Thr Ile Arg Lys Leu Gln Phe Ala Thr Arg  
20 25 30

Arg His Leu Met Ser Ile His Glu Met Gly Gly Ile Arg Asn Ala Asn  
35 40 45

Arg Ile Leu Lys Asp Leu Ser Ile Tyr Thr Ser Lys Val Val Tyr Asn  
50 55 60

Lys Glu His Val Tyr Tyr Leu Asn Gln Ser Gly His Lys Leu Phe Gly  
65 70 75 80

Glu Gly Lys Val Val His His Gly Lys Val Thr His Ala Leu Leu Arg  
85 90 95

Asn Glu Ala Trp Leu Asn Leu Tyr Cys Pro Asp Asp Trp Gln Val Glu  
Page 98

Thr Glu Ile Lys Tyr Ile Lys Asp Asn Lys Lys Lys Lys Ile Ile Pro  
115 120 125

Asp Val Lys Phe Arg Asp Glu Asp Arg Ile Leu His Ala Val Glu Ile  
130 135 140

Asp Arg Thr Gln Lys Met Ile Val Asn Asp Glu Lys Leu Lys Lys Tyr  
145 150 155 160

Glu Glu Leu Thr Gln Ile Tyr Lys Gln Lys His Asn Gly Lys Val Pro  
165 170 175

Val Ile His Phe Phe Thr Ile Thr Lys Tyr Arg Glu Lys Lys Leu Glu  
180 185 190

Glu Leu Ala Asn Lys Tyr Asn Val Phe Val Lys Val Tyr Val Ile Ala  
195 200 205

Thr Thr  
210

<210> 49  
<211> 78  
<212> PRT  
<213> Bacillus anthracis

<400> 49

Met Lys Phe Thr Leu Gly Asn Ser Leu Asp Glu Leu Gly Ile Thr Lys  
1 5 10 15

Asn Lys Leu Ser Thr Glu Ser Gln Val Arg Tyr Asn Thr Ile Ser Asp  
20 25 30

Leu Val Asn Gly Asn Ala Asn Ala Val Arg Phe Asp Ser Leu Glu Ala  
35 40 45

Ile Ile Asp Ala Leu Asn Ala Ile Ala Ala Glu Lys Gly Ile Asn Lys  
50 55 60

Ile Tyr Lys Ile Asp Asp Val Ile Gln Tyr Ile Lys Lys Ser  
65 70 75

<210> 50  
<211> 78  
<212> PRT  
<213> Bacillus anthracis

<400> 50

Met Lys Phe Thr Leu Gly Asn Ser Leu Asp Glu Leu Gly Ile Thr Lys  
1 5 10 15

Asn Lys Leu Ser Thr Glu Ser Gln Val Arg Tyr Asn Thr Ile Ser Asp  
20 25 30

Leu Val Asn Gly Asn Ala Asn Ala Val Arg Phe Asp Ser Leu Glu Ala  
35 40 45

Ile Ile Asp Ala Leu Asn Ala Ile Ala Ala Glu Lys Gly Ile Asn Lys  
50 55 60

Ile Tyr Lys Ile Asp Asp Val Ile Gln Tyr Ile Lys Lys Ser  
65 70 75

<210> 51  
<211> 75  
<212> PRT  
<213> Bacillus anthracis

<400> 51

Met Ala Phe Lys Ala Ser Met Ile Ala Ser Ser Glu Ser Lys Arg Thr  
1 5 10 15

Ala Leu Ala Leu Pro Phe Thr Lys Ser Leu Ile Val Leu Tyr Leu Thr  
20 25 30

Trp Asp Ser Val Asp Asn Leu Phe Leu Val Ile Pro Asn Ser Ser Lys  
35 40 45

Glu Phe Pro Ser Val Asn Phe Ile Leu Phe Ser Ser Ala Ala Leu Val  
50 55 60

Ile Leu Tyr Ser Phe Tyr Asn Ile Asn Arg Asn  
65 70 75

<210> 52  
<211> 78  
<212> PRT  
<213> Bacillus anthracis

<400> 52

Met Lys Phe Thr Leu Gly Asn Ser Leu Asp Glu Leu Gly Ile Thr Lys  
1 5 10 15

Asn Lys Leu Ser Thr Glu Ser Gln Val Arg Tyr Asn Thr Ile Ser Asp  
20 25 30

Leu Val Asn Gly Asn Ala Asn Ala Val Arg Phe Asp Ser Leu Glu Ala  
35 40 45

Ile Ile Asp Ala Leu Asn Ala Ile Ala Ala Glu Lys Gly Ile Asn Lys  
50 55 60

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Ile Tyr Lys Ile Asp Asp Val Ile Gln Tyr Ile Lys Lys Ser  
65 70 75

<210> 53  
<211> 287  
<212> PRT  
<213> Bacillus anthracis

<400> 53

Met Leu Ser Ser Ala Asn Tyr Thr Gln Tyr Lys Lys Leu Gln Ser Phe  
1 5 10 15

Arg Ser Val Glu Glu Met Asn Glu Ala Ile Cys Ser Phe Leu Tyr Lys  
20 25 30

His Thr His Glu Leu Ser Glu Ser Ala Ile Lys Val Leu Lys Phe Leu  
35 40 45

Ala Arg His Ser Cys Lys Ile Pro Gly Val Ser Phe Leu Lys Val Gly  
50 55 60

Thr Ile Ala Glu Ala Leu Asn Ile Ser Asp Arg Thr Val Arg Arg Val  
65 70 75 80

Leu Lys Val Leu Glu Asp Phe Glu Val Val Thr Arg His Lys Thr Ile  
85 90 95

Arg Thr Glu Gly Lys Leu Arg Gly Gly Asn Gly His Asn Val Tyr Val  
100 105 110

Leu Leu Lys Lys Tyr Ser Val Thr Pro Asn Val Leu Pro Lys Met Ser  
115 120 125

Gln Arg Gln Asp Glu Glu Asn Leu Thr Glu Ser Lys Val Ser Asp Thr  
130 135 140

Lys Thr Asp Lys Glu Ala Lys Leu Ser Glu Ser His Pro Leu Glu Glu  
145 150 155 160

Leu Lys Ser Glu Leu Asn Val Lys Glu Thr Ser Ala Arg Glu Ser Lys  
165 170 175

Glu Ile Glu Leu Glu Asp Leu Asp Glu Thr Phe Thr Pro Glu Asn Val  
180 185 190

Pro Ser Gln Phe Arg Asp Val Val Ala Pro Phe Phe Lys Ser Ala Asp  
195 200 205

Lys Ile Tyr Lys Leu Tyr His Arg Val Leu Ile Ala Tyr Lys Arg Ser  
210 215 220

PCT-US2005-009928\_Sequence Listing.txt as filed 10-3-05 in PCT

Lys Ile Asp Lys Pro Ile Glu Gln Val Ile Asn Gln Ala Ile Gln Ala  
225 230 235 240

Phe Lys Glu Thr Val Phe Ala Glu Lys Ala Asn Lys Ile Arg Ser Thr  
245 250 255

Phe Glu Gly Tyr Phe Tyr Arg Ile Val Glu Ser Lys Phe Val Met Glu  
260 265 270

Arg Arg Lys Glu Cys Arg Gly Leu Leu Phe Asp Trp Leu Asn Glu  
275 280 285

<210> 54  
<211> 287  
<212> PRT  
<213> Bacillus anthracis

<400> 54

Met Leu Ser Ser Ala Asn Tyr Thr Gln Tyr Lys Lys Leu Gln Ser Phe  
1 5 10 15

Arg Ser Val Glu Glu Met Asn Glu Ala Ile Cys Ser Phe Leu Tyr Lys  
20 25 30

His Thr His Glu Leu Ser Glu Ser Ala Ile Lys Val Leu Lys Phe Leu  
35 40 45

Ala Arg His Ser Cys Lys Ile Pro Gly Val Ser Phe Leu Lys Val Gly  
50 55 60

Thr Ile Ala Glu Ala Leu Asn Ile Ser Asp Arg Thr Val Arg Arg Val  
65 70 75 80

Leu Lys Val Leu Glu Asp Phe Glu Val Val Thr Arg His Lys Thr Ile  
85 90 95

Arg Thr Glu Gly Lys Leu Arg Gly Gly Asn Gly His Asn Val Tyr Val  
100 105 110

Leu Leu Lys Lys Tyr Ser Val Thr Pro Asn Val Leu Pro Lys Met Ser  
115 120 125

Gln Arg Gln Asp Glu Glu Asn Leu Thr Glu Ser Lys Val Ser Asp Thr  
130 135 140

Lys Thr Asp Lys Glu Ala Lys Leu Ser Glu Ser His Pro Leu Glu Glu  
145 150 155 160

Leu Lys Ser Glu Leu Asn Val Lys Glu Thr Ser Ala Arg Glu Ser Lys  
165 170 175

PCT-US2005-009928\_Sequence Listing.txt as filed 10-3-05 in PCT

Glu Ile Glu Leu Glu Asp Leu Asp Glu Thr Phe Thr Pro Glu Asn Val  
180 185 190

Pro Ser Gln Phe Arg Asp Val Val Ala Pro Phe Phe Lys Ser Ala Asp  
195 200 205

Lys Ile Tyr Lys Leu Tyr His Arg Val Leu Ile Ala Tyr Lys Arg Ser  
210 215 220

Lys Ile Asp Lys Pro Ile Glu Gln Val Ile Asn Gln Ala Ile Gln Ala  
225 230 235 240

Phe Lys Glu Thr Val Phe Ala Glu Lys Ala Asn Lys Ile Arg Ser Thr  
245 250 255

Phe Glu Gly Tyr Phe Tyr Arg Ile Val Glu Ser Lys Phe Val Met Glu  
260 265 270

Arg Arg Lys Glu Cys Arg Gly Leu Leu Phe Asp Trp Leu Asn Glu  
275 280 285

<210> 55  
<211> 481  
<212> PRT  
<213> Bacillus anthracis

<400> 55

Leu Lys Tyr Ala Val Tyr Val Arg Val Ser Thr Asp Arg Asp Glu Gln  
1 5 10 15

Val Ser Ser Val Glu Asn Gln Ile Asp Ile Cys Arg Tyr Trp Leu Glu  
20 25 30

Lys Asn Gly Tyr Glu Trp Asp Pro Asn Ala Val Tyr Phe Asp Asp Gly  
35 40 45

Ile Ser Gly Thr Ala Trp Leu Glu Arg His Ala Met Gln Leu Ile Leu  
50 55 60

Glu Lys Ala Arg Arg Asn Glu Leu Asp Thr Val Val Phe Lys Ser Ile  
65 70 75 80

His Arg Leu Ala Arg Asp Leu Arg Asp Ala Leu Glu Ile Lys Glu Ile  
85 90 95

Leu Ile Gly His Gly Ile Arg Leu Val Thr Ile Glu Glu Asn Tyr Asp  
100 105 110

Ser Leu Tyr Glu Gly Gly Asn Asp Ile Lys Phe Glu Met Phe Ala Met  
115 120 125



PCT-US2005-009928\_Sequence Listing.txt as filed 10-3-05 in PCT

Phe	Ala	Ala	Gln	Leu	Pro	Lys	Thr	Ile	Ser	Val	Ser	Val	Ser	Ala	Ala
	130					135					140				
Met	Gln	Ala	Lys	Ala	Arg	Arg	Gly	Glu	Phe	Ile	Gly	Lys	Pro	Gly	Leu
145					150					155					160
Gly	Tyr	Asp	Val	Ile	Asp	Lys	Lys	Leu	Val	Ile	Asn	Glu	Lys	Glu	Ala
				165					170					175	
Glu	Ile	Val	Arg	Glu	Ile	Phe	Asp	Leu	Ser	Tyr	Lys	Gly	Tyr	Gly	Phe
			180					185					190		
Lys	Lys	Ile	Ala	Asn	Ile	Leu	Asn	Asp	Lys	Gly	Thr	Tyr	Thr	Lys	Phe
		195					200					205			
Gly	Gln	Leu	Trp	Ser	His	Thr	Thr	Val	Gly	Lys	Ile	Leu	Lys	Asn	Gln
	210					215					220				
Thr	Tyr	Lys	Gly	Asn	Leu	Val	Leu	Asn	Ser	Tyr	Lys	Thr	Val	Lys	Val
225					230					235					240
Asp	Gly	Lys	Lys	Lys	Arg	Val	Tyr	Thr	Pro	Lys	Glu	Arg	Leu	Thr	Ile
				245					250					255	
Ile	Glu	Asp	His	Tyr	Pro	Thr	Ile	Val	Ser	Lys	Glu	Leu	Trp	Asn	Ala
			260					265					270		
Val	Asn	Ser	Asp	Arg	Ala	Ser	Lys	Lys	Lys	Thr	Lys	Gln	Asp	Thr	Arg
		275					280					285			
Asn	Glu	Phe	Arg	Gly	Met	Met	Phe	Cys	Lys	His	Cys	Gly	Glu	Pro	Ile
	290					295					300				
Thr	Ala	Lys	Tyr	Ser	Gly	Arg	Tyr	Ala	Lys	Gly	Ser	Lys	Lys	Glu	Trp
305					310					315					320
Val	Tyr	Met	Lys	Cys	Ser	Asn	Tyr	Ile	Arg	Phe	Asn	Arg	Cys	Val	Asn
				325					330					335	
Phe	Asp	Pro	Ala	His	Tyr	Asp	Asp	Ile	Arg	Glu	Ala	Ile	Ile	Tyr	Gly
			340					345					350		
Leu	Lys	Gln	Gln	Glu	Lys	Glu	Leu	Glu	Ile	His	Phe	Asn	Pro	Lys	Met
		355					360					365			
His	Gln	Lys	Arg	Asn	Asp	Lys	Ser	Thr	Glu	Ile	Lys	Lys	Gln	Ile	Lys
	370					375					380				
Leu	Leu	Lys	Val	Lys	Lys	Glu	Lys	Leu	Ile	Asp	Leu	Tyr	Val	Glu	Gly
385					390					395					400

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Leu Ile Asp Lys Glu Met Phe Ser Lys Arg Asp Leu Asn Phe Glu Asn  
405 410 415

Glu Ile Lys Glu Gln Glu Leu Ala Leu Leu Lys Leu Thr Asp Gln Asn  
420 425 430

Lys Arg Asn Lys Glu Glu Lys Lys Ile Lys Glu Ala Phe Ser Met Leu  
435 440 445

Asp Glu Glu Lys Asp Met His Glu Val Phe Lys Thr Leu Ile Lys Lys  
450 455 460

Ile Thr Leu Ser Lys Asp Lys Tyr Ile Asp Ile Glu Tyr Thr Phe Ser  
465 470 475 480

Leu

<210> 56  
<211> 481  
<212> PRT  
<213> Bacillus anthracis

<400> 56

Leu Lys Tyr Ala Val Tyr Val Arg Val Ser Thr Asp Arg Asp Glu Gln  
1 5 10 15

Val Ser Ser Val Glu Asn Gln Ile Asp Ile Cys Arg Tyr Trp Leu Glu  
20 25 30

Lys Asn Gly Tyr Glu Trp Asp Pro Asn Ala Val Tyr Phe Asp Asp Gly  
35 40 45

Ile Ser Gly Thr Ala Trp Leu Glu Arg His Ala Met Gln Leu Ile Leu  
50 55 60

Glu Lys Ala Arg Arg Asn Glu Leu Asp Thr Val Val Phe Lys Ser Ile  
65 70 75 80

His Arg Leu Ala Arg Asp Leu Arg Asp Ala Leu Glu Ile Lys Glu Ile  
85 90 95

Leu Ile Gly His Gly Ile Arg Leu Val Thr Ile Glu Glu Asn Tyr Asp  
100 105 110

Ser Leu Tyr Glu Gly Gly Asn Asp Ile Lys Phe Glu Met Phe Ala Met  
115 120 125

Phe Ala Ala Gln Leu Pro Lys Thr Ile Ser Val Ser Val Ser Ala Ala  
130 135 140

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Met	Gln	Ala	Lys	Ala	Arg	Arg	Gly	Glu	Phe	Ile	Gly	Lys	Pro	Gly	Leu
145					150					155					160
Gly	Tyr	Asp	Val	Ile	Asp	Lys	Lys	Leu	Val	Ile	Asn	Glu	Lys	Glu	Ala
				165					170					175	
Glu	Ile	Val	Arg	Glu	Ile	Phe	Asp	Leu	Ser	Tyr	Lys	Gly	Tyr	Gly	Phe
			180					185					190		
Lys	Lys	Ile	Ala	Asn	Ile	Leu	Asn	Asp	Lys	Gly	Thr	Tyr	Thr	Lys	Phe
		195					200					205			
Gly	Gln	Leu	Trp	Ser	His	Thr	Thr	Val	Gly	Lys	Ile	Leu	Lys	Asn	Gln
	210					215					220				
Thr	Tyr	Lys	Gly	Asn	Leu	Val	Leu	Asn	Ser	Tyr	Lys	Thr	Val	Lys	Val
225					230					235					240
Asp	Gly	Lys	Lys	Lys	Arg	Val	Tyr	Thr	Pro	Lys	Glu	Arg	Leu	Thr	Ile
				245					250					255	
Ile	Glu	Asp	His	Tyr	Pro	Thr	Ile	Val	Ser	Lys	Glu	Leu	Trp	Asn	Ala
			260					265					270		
Val	Asn	Ser	Asp	Arg	Ala	Ser	Lys	Lys	Lys	Thr	Lys	Gln	Asp	Thr	Arg
		275					280					285			
Asn	Glu	Phe	Arg	Gly	Met	Met	Phe	Cys	Lys	His	Cys	Gly	Glu	Pro	Ile
	290					295					300				
Thr	Ala	Lys	Tyr	Ser	Gly	Arg	Tyr	Ala	Lys	Gly	Ser	Lys	Lys	Glu	Trp
305					310					315					320
Val	Tyr	Met	Lys	Cys	Ser	Asn	Tyr	Ile	Arg	Phe	Asn	Arg	Cys	Val	Asn
				325					330					335	
Phe	Asp	Pro	Ala	His	Tyr	Asp	Asp	Ile	Arg	Glu	Ala	Ile	Ile	Tyr	Gly
			340					345					350		
Leu	Lys	Gln	Gln	Glu	Lys	Glu	Leu	Glu	Ile	His	Phe	Asn	Pro	Lys	Met
		355					360					365			
His	Gln	Lys	Arg	Asn	Asp	Lys	Ser	Thr	Glu	Ile	Lys	Lys	Gln	Ile	Lys
	370					375					380				
Leu	Leu	Lys	Val	Lys	Lys	Glu	Lys	Leu	Ile	Asp	Leu	Tyr	Val	Glu	Gly
385					390					395					400
Leu	Ile	Asp	Lys	Glu	Met	Phe	Ser	Lys	Arg	Asp	Leu	Asn	Phe	Glu	Asn

Glu Ile Lys Glu Gln Glu Leu Ala Leu Leu Lys Leu Thr Asp Gln Asn  
420 425 430

Lys Arg Asn Lys Glu Glu Lys Lys Ile Lys Glu Ala Phe Ser Met Leu  
435 440 445

Asp Glu Glu Lys Asp Met His Glu Val Phe Lys Thr Leu Ile Lys Lys  
450 455 460

Ile Thr Leu Ser Lys Asp Lys Tyr Ile Asp Ile Glu Tyr Thr Phe Ser  
465 470 475 480

Leu

<210> 57  
<211> 42  
<212> PRT  
<213> Bacillus anthracis

<400> 57

Val Ile Ile Val Glu Phe Lys Asp Arg Leu Arg Gln Leu Arg Arg Glu  
1 5 10 15

Arg Asn Leu Thr Gln His Asp Leu Gly Gln Ala Ile Gly Val Thr Ala  
20 25 30

Gly Ser Ile Thr Val Thr Asn Asn Gln Leu  
35 40

<210> 58  
<211> 444  
<212> PRT  
<213> Bacillus anthracis

<400> 58

Met Arg Ile Ala Leu Tyr Arg Thr His Ala Leu Ile Asn Val Ile Lys  
1 5 10 15

Tyr Ser Val Asn Ile Met Glu Lys Val Leu Leu Ile Glu Met Lys Gly  
20 25 30

Val Ser Tyr Leu Lys Phe His Glu Lys Ile Met Gly Met Ile Glu Asp  
35 40 45

Arg Asp Asp Leu Thr Ala Thr Ser Val Ala Cys Lys Ile Gly Val Ser  
50 55 60

Lys Gln Tyr Met Ser Lys Phe Lys Arg Gln Gly Thr Ile Gly Phe Ser  
65 70 75 80

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Gln	Leu	Leu	Lys	Leu	Ala	Pro	Ile	Leu	Ser	Val	Glu	Gly	Lys	Lys	Ala
			85						90					95	
Lys	Gln	Thr	Met	Ser	Asp	Trp	Cys	Leu	Glu	Leu	Asp	Thr	Thr	Glu	Ser
			100					105					110		
Ile	Lys	Gln	Ser	Phe	Glu	Tyr	Ala	Cys	Leu	Thr	Arg	Asn	Thr	Ile	Leu
		115					120					125			
Leu	Lys	Gln	Leu	Ile	Gln	Lys	His	Ser	Lys	Glu	Thr	Gly	Thr	Ile	Arg
	130					135					140				
Glu	Tyr	Val	Glu	Val	Tyr	Thr	Ile	Leu	Phe	Lys	Tyr	Ile	Lys	Asn	Ile
145					150					155					160
Ile	Lys	Gly	Ser	Glu	Ile	Thr	Lys	Glu	Leu	Lys	Lys	Ile	Gly	Ala	Ile
				165					170					175	
Lys	Asp	Lys	Val	Leu	Glu	Ile	Leu	Thr	Lys	Ile	Met	Glu	Cys	Tyr	Glu
			180					185					190		
Tyr	Tyr	His	Leu	Lys	Lys	Phe	Asn	Leu	Met	Leu	Glu	Thr	Ala	Glu	Thr
		195					200					205			
Ile	Asp	Ser	Leu	Val	Arg	Glu	Ile	Glu	Gly	Glu	Arg	Lys	Ser	Phe	Ile
	210					215					220				
Lys	Glu	Cys	Tyr	Asn	Tyr	Arg	Ile	Ala	Glu	Leu	Phe	Ala	Pro	Ile	Phe
225					230					235					240
Leu	Gln	Lys	Asn	Asn	Val	Asp	Leu	Ala	Arg	Lys	Tyr	Ala	His	Phe	Leu
				245					250					255	
Ile	His	Ala	Asn	Val	Cys	Thr	Lys	Thr	Val	Ser	Asp	Ala	Tyr	Tyr	Ile
			260					265					270		
Leu	Gly	Met	Ser	Asn	Val	Leu	Glu	Ser	Lys	Glu	Gln	Cys	Leu	Phe	Asn
		275					280					285			
Leu	Lys	Lys	Ser	Tyr	Leu	Leu	Ser	Lys	Glu	Ile	Arg	Asp	Ala	Asp	Ile
	290					295					300				
Glu	Gln	Glu	Ala	Arg	Tyr	Asn	Leu	Asp	Val	Ala	Lys	Ile	Tyr	Phe	Gly
305					310					315					320
Val	Lys	Leu	Asp	Glu	Asp	Ala	Asp	Ser	Arg	Leu	Leu	Leu	Tyr	Gln	Lys
				325					330					335	
Asn	Pro	Thr	Cys	Glu	Leu	Ser	Ile	Ile	Ala	Leu	Gln	Asp	Ile	Ile	Arg

Asp Arg Gly Asp Lys Asp Phe Leu Asn Tyr Phe Ile Ala Cys Ser Ser  
355 360 365

Asp Glu Ile Glu Cys Leu Tyr Asp Leu Phe Tyr Gln Tyr Phe Tyr Gln  
370 375 380

Ala Asn Tyr Leu Phe Ser Ala Ile Val Ala Lys Glu Leu Cys Asn Arg  
385 390 395 400

Gly Asp Lys Ser Leu Leu Thr Gln Ser Met Val Asn Leu Gly Asn Glu  
405 410 415

Lys Gln Lys Gly Val Val Asp Ile Glu Glu Ile Ser Ile Ser Ser Leu  
420 425 430

Tyr Ile Ile Asn Gly Ser Asn Ser Gly Ile Val Val  
435 440

<210> 59  
<211> 75  
<212> PRT  
<213> Bacillus anthracis

<400> 59

Met Lys Val Ile Lys Asp Glu Thr Lys Leu Lys Ala Ala Phe Lys Lys  
1 5 10 15

Ser Gly Tyr Lys Tyr Gln Glu Leu Ala Asp Glu Leu Glu Ile Ser Cys  
20 25 30

Ser Tyr Cys Tyr Lys Leu Ile Asn Asn His Asn Tyr Lys Lys Lys Ile  
35 40 45

Ser Tyr Asn Leu Ala Ser Arg Met Ala His Val Leu Asn Ala Ser Val  
50 55 60

Val Asp Leu Phe Glu Glu Gln Val Asp Phe Phe  
65 70 75

<210> 60  
<211> 118  
<212> PRT  
<213> Bacillus anthracis

<400> 60

Val Ile Ile Val Glu Phe Lys Asp Arg Leu Arg Gln Leu Arg Arg Glu  
1 5 10 15

Arg Asn Leu Thr Gln His Asp Leu Gly Gln Ala Ile Gly Val Thr Ala  
20 25 30

Gly Ser Val Ser Lys Phe Glu Thr Gly Phe Lys Pro Ala Ser Arg Glu  
35 40 45

Thr Val Glu Arg Ala Ala Asp Phe Leu Gly Val Pro Val Asp Tyr Leu  
50 55 60

Leu Gly Arg Ser Asp Ser Arg Glu Leu Asp Ala Asp Met Asn Gln Lys  
65 70 75 80

Tyr Leu His Ile Lys Asn Arg Leu Glu Gln Leu Pro Glu Glu His Gln  
85 90 95

Glu Ile Val Leu Gln Asn Met Leu Thr Met Met Glu Ser Leu Glu Lys  
100 105 110

Leu Lys Ser Thr Ser Lys  
115

<210> 61  
<211> 61  
<212> PRT  
<213> Bacillus anthracis

<400> 61

Met Arg Glu His Arg Gly Glu Arg Ala Met Ser Glu Ile Tyr Tyr Lys  
1 5 10 15

Gly Phe Ile Ile Lys Glu Thr Tyr Gly Glu Arg Asn Ile Glu Glu Val  
20 25 30

Phe Lys Glu Ala Tyr Glu Ser Phe Tyr Gly Val Glu Val Lys Val Val  
35 40 45

Lys Lys Glu Leu Gly Thr Lys Arg Asn Ser Ala Ala Ser  
50 55 60

<210> 62  
<211> 75  
<212> PRT  
<213> Bacillus anthracis

<400> 62

Met Lys Val Ile Lys Asp Glu Thr Lys Leu Lys Ala Ala Phe Lys Lys  
1 5 10 15

Ser Gly Tyr Lys Tyr Gln Glu Leu Ala Asp Glu Leu Glu Ile Ser Cys  
20 25 30

Ser Tyr Cys Tyr Lys Leu Ile Asn Asn His Asn Tyr Lys Lys Lys Ile  
35 40 45



PCT-US2005-009928\_Sequence Listing.txt as filed 10-3-05 in PCT

Ser Tyr Asn Leu Ala Ser Arg Met Ala His Val Leu Asn Ala Ser Val  
50 55 60

Val Asp Leu Phe Glu Glu Gln Val Asp Phe Phe  
65 70 75

<210> 63  
<211> 271  
<212> PRT  
<213> Bacillus anthracis

<400> 63

Met Asp Gln Leu Thr Val Ala Ser Glu Leu Arg Leu Leu Gly Arg Arg  
1 5 10 15

Lys Val Ala Gly Tyr Glu Phe Thr Gly Ile Glu Gly Gly Phe Gly Glu  
20 25 30

Gly Lys Lys Ala Met Leu Val Leu Asp Ile Ala Thr Ile His Asn Gln  
35 40 45

Pro Leu Lys Glu Ile Asn Arg Arg Ile Asn Asp Asn Arg Ile Arg Phe  
50 55 60

Lys Asp Gly Val Asp Ile Val Asp Leu Lys Ser Gly Gly Phe Asn Pro  
65 70 75 80

Pro Gln Leu Leu Asn Leu Gly Phe Ser Asn Met Gln Ile Ala Lys Ser  
85 90 95

Asn Asn Ile Tyr Leu Leu Ser Glu Arg Gly Tyr Ala Lys Leu Leu Lys  
100 105 110

Ile Leu Glu Asp Asp Lys Ala Trp Glu Leu Tyr Asp Ile Leu Val Asp  
115 120 125

Glu Tyr Phe Asn Met Arg Glu Lys Asn Gln Val Ala Thr Asp Pro Met  
130 135 140

Ser Ile Leu Lys Leu Thr Phe Glu Ala Leu Glu Gly Gln Gln Gln Ala  
145 150 155 160

Ile Glu Glu Ile Lys Ser Asp Val Gln Asp Leu Arg Glu Asn Thr Pro  
165 170 175

Leu Phe Ala Ile Glu Cys Asp Glu Ile Ser Thr Ala Val Lys Arg Gln  
180 185 190

Gly Val Ile Leu Leu Gly Gly Lys Gln Ser Asn Ala Tyr Arg Asn Arg  
195 200 205

PCT-US2005-009928\_sequence Listing.txt as filed 10-3-05 in PCT

Gly Leu Arg Gly Lys Val Tyr Arg Asp Ile Tyr Asn Gln Leu Tyr Arg  
210 215 220

Glu Phe Gly Val Lys Ser His Lys Ala Ile Lys Arg Cys His Leu Asn  
225 230 235 240

Val Ala Val Lys Ile Val Glu Glu Tyr Thr Leu Pro Ile Val Leu Ser  
245 250 255

Glu Glu Ile Ser Phe Val Asn Ala Gln Met Asp Phe Thr Glu Met  
260 265 270

<210> 64  
<211> 61  
<212> PRT  
<213> Bacillus anthracis

<400> 64

Met Arg Glu His Arg Gly Glu Arg Ala Met Ser Glu Ile Tyr Tyr Lys  
1 5 10 15

Gly Phe Ile Ile Lys Glu Thr Tyr Gly Glu Arg Asn Ile Glu Glu Val  
20 25 30

Phe Lys Glu Ala Tyr Glu Ser Phe Tyr Gly Val Glu Val Lys Val Val  
35 40 45

Lys Lys Glu Leu Gly Thr Lys Arg Asn Ser Ala Ala Ser  
50 55 60

<210> 65  
<211> 217  
<212> PRT  
<213> Bacillus anthracis

<400> 65

Met Asp Gln Leu Arg Val Ile Glu Gly Glu Lys Val Asp Lys Pro Asp  
1 5 10 15

Tyr Val Glu Ile Tyr Leu Gly Ala Phe Met Asn Ala Val Asn Glu Leu  
20 25 30

Lys Lys Gln Asp Glu Glu Thr Arg Ser Leu Ser Lys Asp Thr Tyr Lys  
35 40 45

Lys Ala Ile Phe Tyr Gly Val Arg Tyr Ile Ser Ile Ser Lys Asn Asp  
50 55 60

Ser Leu Asn Tyr Asp Tyr Leu Met Asn Arg Phe Leu Leu Ile Ser Tyr  
65 70 75 80

PCT-US2005-009928\_Sequence Listing.txt as filed 10-3-05 in PCT  
Leu Glu Asn Leu Met Lys Val Leu Thr Pro Arg Asp Phe Met Thr Ile  
85 90 95

Phe Pro Ile Asp Lys Asn Tyr Asp Gly Ala Arg Tyr Glu Met Lys Asp  
100 105 110

Tyr Phe Phe Thr Met Asn Glu Ile Lys Lys Ile Gly Met Asp Thr Pro  
115 120 125

Ile Gly Glu Lys Ile Met Glu Phe Leu Trp Asp Tyr Gln Asn Phe Lys  
130 135 140

Asp Ile Thr Leu Phe Asn Leu Ala Ser Val Ser Ile Leu Asn Lys Leu  
145 150 155 160

Gln Lys Met Gln Gly Lys Lys Thr Leu Thr Glu Glu Phe Ala Glu Arg  
165 170 175

Leu Gly Ile Asp Thr Tyr Thr Lys His Lys Glu Lys Gly Gly Lys Glu  
180 185 190

Tyr Ile Thr Asn Asp Arg Thr Gly Glu Ile Gln Glu Val Lys Lys Ser  
195 200 205

Arg Pro Arg Tyr Leu Lys Pro Val Gln  
210 215

<210> 66  
<211> 271  
<212> PRT  
<213> Bacillus anthracis

<400> 66

Met Asp Gln Leu Thr Val Ala Ser Glu Leu Arg Leu Leu Gly Arg Arg  
1 5 10 15

Lys Val Ala Gly Tyr Glu Phe Thr Gly Ile Glu Gly Gly Phe Gly Glu  
20 25 30

Gly Lys Lys Ala Met Leu Val Leu Asp Ile Ala Thr Ile His Asn Gln  
35 40 45

Pro Leu Lys Glu Ile Asn Arg Arg Ile Asn Asp Asn Arg Ile Arg Phe  
50 55 60

Lys Asp Gly Val Asp Ile Val Asp Leu Lys Ser Gly Gly Phe Asn Pro  
65 70 75 80

Pro Gln Leu Leu Asn Leu Gly Phe Ser Asn Met Gln Ile Ala Lys Ser  
85 90 95

Asn Asn Ile Tyr Leu Leu Ser Glu Arg Gly Tyr Ala Lys Leu Leu Lys  
100 105 110

Ile Leu Glu Asp Asp Lys Ala Trp Glu Leu Tyr Asp Ile Leu Val Asp  
115 120 125

Glu Tyr Phe Asn Met Arg Glu Lys Asn Gln Val Ala Thr Asp Pro Met  
130 135 140

Ser Ile Leu Lys Leu Thr Phe Glu Ala Leu Glu Gly Gln Gln Gln Ala  
145 150 155 160

Ile Glu Glu Ile Lys Ser Asp Val Gln Asp Leu Arg Glu Asn Thr Pro  
165 170 175

Leu Phe Ala Ile Glu Cys Asp Glu Ile Ser Thr Ala Val Lys Arg Gln  
180 185 190

Gly Val Ile Leu Leu Gly Gly Lys Gln Ser Asn Ala Tyr Arg Asn Arg  
195 200 205

Gly Leu Arg Gly Lys Val Tyr Arg Asp Ile Tyr Asn Gln Leu Tyr Arg  
210 215 220

Glu Phe Gly Val Lys Ser His Lys Ala Ile Lys Arg Cys His Leu Asn  
225 230 235 240

Val Ala Val Lys Ile Val Glu Glu Tyr Thr Leu Pro Ile Val Leu Ser  
245 250 255

Glu Glu Ile Ser Phe Val Asn Ala Gln Met Asp Phe Thr Glu Met  
260 265 270

<210> 67  
<211> 315  
<212> PRT  
<213> Bacillus anthracis

<400> 67

Met Ala Leu Phe Arg Lys Val His Thr Glu Phe Trp Thr Asp Val Lys  
1 5 10 15

Val Ser Glu Asp Met Thr Pro Glu Asp Lys Leu Phe Met Val Tyr Leu  
20 25 30

Leu Thr Asn Pro His Thr Thr Gln Leu Gly Val Tyr Glu Ile Thr Pro  
35 40 45

Lys Met Ile Ala Phe Glu Ile Gly Leu Ser Ile Glu Ser Ala Arg Ala  
50 55 60

Leu Leu Glu Arg Phe Glu Asn His His Lys Leu Ile Lys Tyr Asn Lys  
65 70 75 80

Leu Thr Arg Glu Ile Ala Ile Lys Asn Trp Gly Lys Tyr Asn Leu Asn  
85 90 95

Arg Gly Gly Lys Pro Ile Glu Asp Cys Leu Lys Arg Glu Ile Asp Lys  
100 105 110

Val Lys Asp Leu Ser Leu Ile Lys Phe Ile Leu Glu His Thr Asp His  
115 120 125

Ala Ala Leu Lys Arg Lys Ile Asn Leu Tyr Ala Gly Phe Asp Asp Thr  
130 135 140

Ser His Asp Thr Leu Ala Ile Arg Asp Gln Glu Glu Glu Lys Glu Gln  
145 150 155 160

Lys Lys Glu Gln Lys Glu Glu Gln Glu Glu Lys Glu Lys Glu Lys Glu  
165 170 175

Lys Gln Lys Glu Glu Glu Lys Glu Pro Glu Glu Glu Lys Thr Arg Ile  
180 185 190

Lys Ser Lys Ala Ser Leu Lys Ser Asp Ala Lys Ser Asn Pro Ile Pro  
195 200 205

Tyr Lys Asp Ile Leu Asp Tyr Leu Asn Glu Lys Ala Asn Lys Asn Phe  
210 215 220

Asn Pro Lys Ala Glu Gly His Arg Lys Leu Ile Arg Ala Arg Trp Asn  
225 230 235 240

Glu Gly Tyr Lys Leu Glu Asp Phe Lys Lys Val Ile Asp Asn Lys Thr  
245 250 255

Thr Gln Trp Phe Gly Lys Lys Ser Phe Asp Gly Lys Pro Leu Asp Gln  
260 265 270

Phe Leu Arg Pro Ser Thr Leu Phe Ala Gln Lys His Phe Asp Asn Tyr  
275 280 285

Leu Asn Glu Thr Val Asn Ile Ser Asn Gln Gln His Gly Asp Gln Ile  
290 295 300

Val Ile Pro Gly Phe Arg Gly Glu Met Pro Phe  
305 310 315

<210> 68  
<211> 217  
<212> PRT

PCT-US2005-009928\_Sequence Listing.txt as filed 10-3-05 in PCT  
<213> Bacillus anthracis

<400> 68

Met Asp Gln Leu Arg Val Ile Glu Gly Glu Lys Val Asp Lys Pro Asp  
1 5 10 15  
Tyr Val Glu Ile Tyr Leu Gly Ala Phe Met Asn Ala Val Asn Glu Leu  
20 25 30  
Lys Lys Gln Asp Glu Glu Thr Arg Ser Leu Ser Lys Asp Thr Tyr Lys  
35 40 45  
Lys Ala Ile Phe Tyr Gly Val Arg Tyr Ile Ser Ile Ser Lys Asn Asp  
50 55 60  
Ser Leu Asn Tyr Asp Tyr Leu Met Asn Arg Phe Leu Leu Ile Ser Tyr  
65 70 75 80  
Leu Glu Asn Leu Met Lys Val Leu Thr Pro Arg Asp Phe Met Thr Ile  
85 90 95  
Phe Pro Ile Asp Lys Asn Tyr Asp Gly Ala Arg Tyr Glu Met Lys Asp  
100 105 110  
Tyr Phe Phe Thr Met Asn Glu Ile Lys Lys Ile Gly Met Asp Thr Pro  
115 120 125  
Ile Gly Glu Lys Ile Met Glu Phe Leu Trp Asp Tyr Gln Asn Phe Lys  
130 135 140  
Asp Ile Thr Leu Phe Asn Leu Ala Ser Val Ser Ile Leu Asn Lys Leu  
145 150 155 160  
Gln Lys Met Gln Gly Lys Lys Thr Leu Thr Glu Glu Phe Ala Glu Arg  
165 170 175  
Leu Gly Ile Asp Thr Tyr Thr Lys His Lys Glu Lys Gly Gly Lys Glu  
180 185 190  
Tyr Ile Thr Asn Asp Arg Thr Gly Glu Ile Gln Glu Val Lys Lys Ser  
195 200 205  
Arg Pro Arg Tyr Leu Lys Pro Val Gln  
210 215

<210> 69  
<211> 303  
<212> PRT  
<213> Bacillus anthracis

<400> 69

PCT-US2005-009928\_Sequence Listing.txt as filed 10-3-05 in PCT

Val Lys Lys Ile Gln Asp Ser Phe Glu Lys Leu Thr Lys Leu Lys Phe  
1 5 10 15

Ala Asp Glu Gln Cys Asp Lys His Thr Phe Asn Lys His Gly Lys Glu  
20 25 30

Val Ile Lys Leu Val Arg Lys Met Ile Asp Asp Ala Gly Thr Val Tyr  
35 40 45

Cys Pro Arg Cys Met Val Glu Glu Gln Asn Ser Val Leu Phe Gln Gln  
50 55 60

Ala Asn Asn His Tyr Lys Lys Ile Asn Arg Glu Arg Lys Lys Asn Val  
65 70 75 80

Leu Phe Gln His Ser Ile Ile Glu Asn Gln Ser Ile Thr Glu Ser Arg  
85 90 95

Leu Ser Thr Tyr Lys Thr Asp Cys Gln Glu Thr Lys Glu Asn Lys Glu  
100 105 110

Lys Ala Ile Lys Ile Leu Glu Arg Ile Lys Asn Gly Glu Phe Leu Asn  
115 120 125

Val Tyr Ile Ala Gly Ile Gln Gly Val Gly Lys Ser His Leu Ala Tyr  
130 135 140

Ala Met Leu Tyr Glu Leu Val Lys His Tyr Trp Val Ile Ser Asp Gly  
145 150 155 160

Glu Lys Leu Asn Asp Glu His Ala Phe Lys Asn Met Lys Ser Cys Leu  
165 170 175

Phe Val Glu Ile Glu Lys Leu Ile Arg Leu Ile Gln His Ser Phe Arg  
180 185 190

Asn Ile Glu Ser Lys Tyr Thr Met Asp Tyr Cys Ile Ser Leu Met Val  
195 200 205

Asp Val Asp Phe Leu Val Ile Asp Asp Leu Gly Ala Glu Ser Gly Ser  
210 215 220

Met Asn Arg Asn Gly Glu Ala Ser Asp Phe Val His Lys Ile Leu Tyr  
225 230 235 240

Gly Val Thr Asn Gly Arg Gln Gly Ala Asn Lys Thr Thr Ile Thr Thr  
245 250 255

Ser Asn Leu Ser Ser Ala Gln Leu Phe Gln Lys Tyr Asp Pro Lys Leu  
260 265 270



PCT-US2005-009928\_Sequence Listing.txt as filed 10-3-05 in PCT

Ala Ser Arg Leu Leu Asn Gly Val Ser Lys Asp Glu Thr Ile Val Phe  
275 280 285

Lys Thr Thr Thr Asp Lys Arg Ile Val Asn Leu Asp Ile Gly Phe  
290 295 300

<210> 70  
<211> 315  
<212> PRT  
<213> Bacillus anthracis

<400> 70

Met Ala Leu Phe Arg Lys Val His Thr Glu Phe Trp Thr Asp Val Lys  
1 5 10 15

Val Ser Glu Asp Met Thr Pro Glu Asp Lys Leu Phe Met Val Tyr Leu  
20 25 30

Leu Thr Asn Pro His Thr Thr Gln Leu Gly Val Tyr Glu Ile Thr Pro  
35 40 45

Lys Met Ile Ala Phe Glu Ile Gly Leu Ser Ile Glu Ser Ala Arg Ala  
50 55 60

Leu Leu Glu Arg Phe Glu Asn His His Lys Leu Ile Lys Tyr Asn Lys  
65 70 75 80

Leu Thr Arg Glu Ile Ala Ile Lys Asn Trp Gly Lys Tyr Asn Leu Asn  
85 90 95

Arg Gly Gly Lys Pro Ile Glu Asp Cys Leu Lys Arg Glu Ile Asp Lys  
100 105 110

Val Lys Asp Leu Ser Leu Ile Lys Phe Ile Leu Glu His Thr Asp His  
115 120 125

Ala Ala Leu Lys Arg Lys Ile Asn Leu Tyr Ala Gly Phe Asp Asp Thr  
130 135 140

Ser His Asp Thr Leu Ala Ile Arg Asp Gln Glu Glu Glu Lys Glu Gln  
145 150 155 160

Lys Lys Glu Gln Lys Glu Glu Gln Glu Glu Lys Glu Lys Glu Lys Glu  
165 170 175

Lys Gln Lys Glu Glu Glu Lys Glu Pro Glu Glu Glu Lys Thr Arg Ile  
180 185 190

Lys Ser Lys Ala Ser Leu Lys Ser Asp Ala Lys Ser Asn Pro Ile Pro  
195 200 205

PCT-US2005-009928\_Sequence Listing.txt as filed 10-3-05 in PCT

Tyr Lys Asp Ile Leu Asp Tyr Leu Asn Glu Lys Ala Asn Lys Asn Phe  
210 215 220

Asn Pro Lys Ala Glu Gly His Arg Lys Leu Ile Arg Ala Arg Trp Asn  
225 230 235 240

Glu Gly Tyr Lys Leu Glu Asp Phe Lys Lys Val Ile Asp Asn Lys Thr  
245 250 255

Thr Gln Trp Phe Gly Lys Lys Ser Phe Asp Gly Lys Pro Leu Asp Gln  
260 265 270

Phe Leu Arg Pro Ser Thr Leu Phe Ala Gln Lys His Phe Asp Asn Tyr  
275 280 285

Leu Asn Glu Thr Val Asn Ile Ser Asn Gln Gln His Gly Asp Gln Ile  
290 295 300

Val Ile Pro Gly Phe Arg Gly Glu Met Pro Phe  
305 310 315

<210> 71  
<211> 77  
<212> PRT  
<213> Bacillus anthracis

<400> 71

Met Lys Glu Val Lys Gly Lys Asn Thr Lys Leu Met Glu Glu Phe Asp  
1 5 10 15

Val Leu Leu Arg Gln Leu Leu Ile Lys Ser Lys Thr Asp Glu Arg Val  
20 25 30

Lys Asn Phe Leu Asp Asp Leu Phe Glu Met Leu Ser Asp Asn Lys Leu  
35 40 45

Gln Ser Asp Ile Asp Phe Lys Thr Ala Leu Asn Lys Leu Arg Glu Lys  
50 55 60

His Phe Pro Lys Phe Asp Lys Gly Glu Ser Lys Asn Asp  
65 70 75

<210> 72  
<211> 303  
<212> PRT  
<213> Bacillus anthracis

<400> 72

Val Lys Lys Ile Gln Asp Ser Phe Glu Lys Leu Thr Lys Leu Lys Phe  
1 5 10 15

Ala Asp Glu Gln Cys Asp Lys His Thr Phe Asn Lys His Gly Lys Glu  
 20 25 30  
 Val Ile Lys Leu Val Arg Lys Met Ile Asp Asp Ala Gly Thr Val Tyr  
 35 40 45  
 Cys Pro Arg Cys Met Val Glu Glu Gln Asn Ser Val Leu Phe Gln Gln  
 50 55 60  
 Ala Asn Asn His Tyr Lys Lys Ile Asn Arg Glu Arg Lys Lys Asn Val  
 65 70 75 80  
 Leu Phe Gln His Ser Ile Ile Glu Asn Gln Ser Ile Thr Glu Ser Arg  
 85 90 95  
 Leu Ser Thr Tyr Lys Thr Asp Cys Gln Glu Thr Lys Glu Asn Lys Glu  
 100 105 110  
 Lys Ala Ile Lys Ile Leu Glu Arg Ile Lys Asn Gly Glu Phe Leu Asn  
 115 120 125  
 Val Tyr Ile Ala Gly Ile Gln Gly Val Gly Lys Ser His Leu Ala Tyr  
 130 135 140  
 Ala Met Leu Tyr Glu Leu Val Lys His Tyr Trp Val Ile Ser Asp Gly  
 145 150 155 160  
 Glu Lys Leu Asn Asp Glu His Ala Phe Lys Asn Met Lys Ser Cys Leu  
 165 170 175  
 Phe Val Glu Ile Glu Lys Leu Ile Arg Leu Ile Gln His Ser Phe Arg  
 180 185 190  
 Asn Ile Glu Ser Lys Tyr Thr Met Asp Tyr Cys Ile Ser Leu Met Val  
 195 200 205  
 Asp Val Asp Phe Leu Val Ile Asp Asp Leu Gly Ala Glu Ser Gly Ser  
 210 215 220  
 Met Asn Arg Asn Gly Glu Ala Ser Asp Phe Val His Lys Ile Leu Tyr  
 225 230 235 240  
 Gly Val Thr Asn Gly Arg Gln Gly Ala Asn Lys Thr Thr Ile Thr Thr  
 245 250 255  
 Ser Asn Leu Ser Ser Ala Gln Leu Phe Gln Lys Tyr Asp Pro Lys Leu  
 260 265 270  
 Ala Ser Arg Leu Leu Asn Gly Val Ser Lys Asp Glu Thr Ile Val Phe  
 275 280 285

Lys Thr Thr Thr Asp Lys Arg Ile Val Asn Leu Asp Ile Gly Phe  
290 295 300

<210> 73  
<211> 248  
<212> PRT  
<213> Bacillus anthracis

<400> 73

Met Thr Lys Glu Lys Gly Gln Ala Lys Glu Val Val Asn Val Arg Gly  
1 5 10 15

Met Ser Asp Asp Glu Phe Ile Glu Lys Tyr Gly Arg Leu Val His His  
20 25 30

Cys Val Trp Lys Arg Tyr Ala Lys Lys Lys Ala Ser Ile Glu Arg Asp  
35 40 45

Thr Gly Leu Asp Ile Glu Asp Leu Thr Gln Phe Gly Met Ile Gly Leu  
50 55 60

Ile Lys Ala Arg Asp Asn Phe Asp Leu Glu Phe Gly Cys Ala Phe Ser  
65 70 75 80

Thr Tyr Ala Val Pro Lys Ile Ile Gly Glu Ile Gly Arg Ala Ile Arg  
85 90 95

Asp Asn Gln Lys Ile Lys Val Gln Arg Thr Val Tyr Gly Val Lys Gly  
100 105 110

Lys Ile Leu Asn Gln Gln Leu Ala Asp Lys Glu Pro Glu Glu Ile Ala  
115 120 125

Asp Ile Leu Asp Glu Ser Val Ser Leu Val Lys Thr Ala Leu Glu Tyr  
130 135 140

Gln Pro Ser Thr Asp Ser Leu Asn Lys Val Val Tyr Ala Ser Gly Ala  
145 150 155 160

Asn Glu Glu Leu Thr Leu Glu Arg Met Ile Glu Asp Thr Lys Thr Glu  
165 170 175

Asp Ile Glu Glu Thr Thr Ile Asn Arg Ala Val Ile Arg Glu Phe Lys  
180 185 190

Ala Ala Leu Pro Pro Lys Glu Tyr Ile Val Leu Asp Met Arg Leu Gln  
195 200 205

Asn Met Thr Gln Gln Asn Ile Ala Asn Gln Met Gly Tyr Ser Gln Val  
210 215 220

PCT-US2005-009928\_Sequence Listing.txt as filed 10-3-05 in PCT

Gln Ile Ser Arg Ile Leu Ala Lys Ile Asn Gln Arg Ala Ala Gln Phe  
225 230 235 240

Gly Lys Glu Gly Gly Leu Gln Asp  
245

<210> 74  
<211> 77  
<212> PRT  
<213> Bacillus anthracis

<400> 74

Met Lys Glu Val Lys Gly Lys Asn Thr Lys Leu Met Glu Glu Phe Asp  
1 5 10 15

Val Leu Leu Arg Gln Leu Leu Ile Lys Ser Lys Thr Asp Glu Arg Val  
20 25 30

Lys Asn Phe Leu Asp Asp Leu Phe Glu Met Leu Ser Asp Asn Lys Leu  
35 40 45

Gln Ser Asp Ile Asp Phe Lys Thr Ala Leu Asn Lys Leu Arg Glu Lys  
50 55 60

His Phe Pro Lys Phe Asp Lys Gly Glu Ser Lys Asn Asp  
65 70 75

<210> 75  
<211> 158  
<212> PRT  
<213> Bacillus anthracis

<400> 75

Leu Ser Val Thr Lys Gly Val Cys Ile Asp Val Asp His Ser Asp Leu  
1 5 10 15

Leu His Glu Lys Val Glu Tyr Phe Leu Phe Pro Ala Lys Pro Ser His  
20 25 30

Tyr Tyr Val Ser Arg Phe Asn Arg Lys Gly Ala His Phe Gly Cys Tyr  
35 40 45

Gln Ala Glu Arg Phe Gln Ile Thr Glu Lys Glu Val Trp Thr Pro Glu  
50 55 60

Pro Gln Pro Asn Leu Pro Glu Leu Asn Thr Ser Leu Phe Tyr Arg Ala  
65 70 75 80

Gln Leu Ile Trp Arg Lys Lys Gly Tyr Lys Asp Lys Pro Leu Lys Asp  
85 90 95

PCT-US2005-009928\_Sequence Listing.txt as filed 10-3-05 in PCT  
Tyr Ile Val Gln Pro Arg Gly Lys His Cys Tyr Phe Trp His Asp Arg  
100 105 110

Glu Arg Lys Lys Phe Cys Gly Cys Phe Pro Leu His Trp Phe Thr Asp  
115 120 125

Phe Val Pro Val Gln Ser His His Ile Glu Glu Lys Thr Arg Glu Glu  
130 135 140

Val Lys Leu Leu Gln Arg Pro Asp Gly Gln Leu Ala Phe Phe  
145 150 155

<210> 76  
<211> 248  
<212> PRT  
<213> Bacillus anthracis  
<400> 76

Met Thr Lys Glu Lys Gly Gln Ala Lys Glu Val Val Asn Val Arg Gly  
1 5 10 15

Met Ser Asp Asp Glu Phe Ile Glu Lys Tyr Gly Arg Leu Val His His  
20 25 30

Cys Val Trp Lys Arg Tyr Ala Lys Lys Lys Ala Ser Ile Glu Arg Asp  
35 40 45

Thr Gly Leu Asp Ile Glu Asp Leu Thr Gln Phe Gly Met Ile Gly Leu  
50 55 60

Ile Lys Ala Arg Asp Asn Phe Asp Leu Glu Phe Gly Cys Ala Phe Ser  
65 70 75 80

Thr Tyr Ala Val Pro Lys Ile Ile Gly Glu Ile Gly Arg Ala Ile Arg  
85 90 95

Asp Asn Gln Lys Ile Lys Val Gln Arg Thr Val Tyr Gly Val Lys Gly  
100 105 110

Lys Ile Leu Asn Gln Gln Leu Ala Asp Lys Glu Pro Glu Glu Ile Ala  
115 120 125

Asp Ile Leu Asp Glu Ser Val Ser Leu Val Lys Thr Ala Leu Glu Tyr  
130 135 140

Gln Pro Ser Thr Asp Ser Leu Asn Lys Val Val Tyr Ala Ser Gly Ala  
145 150 155 160

Asn Glu Glu Leu Thr Leu Glu Arg Met Ile Glu Asp Thr Lys Thr Glu  
165 170 175

PCT-US2005-009928\_Sequence Listing.txt as filed 10-3-05 in PCT  
Asp Ile Glu Glu Thr Thr Ile Asn Arg Ala Val Ile Arg Glu Phe Lys  
180 185 190

Ala Ala Leu Pro Pro Lys Glu Tyr Ile Val Leu Asp Met Arg Leu Gln  
195 200 205

Asn Met Thr Gln Gln Asn Ile Ala Asn Gln Met Gly Tyr Ser Gln Val  
210 215 220

Gln Ile Ser Arg Ile Leu Ala Lys Ile Asn Gln Arg Ala Ala Gln Phe  
225 230 235 240

Gly Lys Glu Gly Gly Leu Gln Asp  
245

<210> 77  
<211> 180  
<212> PRT  
<213> Bacillus anthracis

<400> 77

Met Asp Ile Lys Lys Leu Phe Ala Met Gln Asn Ile Leu Asp Lys Arg  
1 5 10 15

Val Leu Glu Ser Lys Asn Leu Ser Arg Gly Glu Val Phe Glu Phe Arg  
20 25 30

Ile Leu Ala Phe Leu Asp Glu Leu Gly Glu Cys Met Lys Glu Trp Arg  
35 40 45

Val Phe Lys Phe Trp Ser Asp Asp Arg Lys Pro Arg Thr Ser Ile Pro  
50 55 60

Thr Gly Glu Ile Ile Val Leu Asp Asp Gly Tyr Glu Val Glu Val Tyr  
65 70 75 80

Lys Asn Pro Leu Leu Glu Glu Tyr Val Asp Gly Leu His Phe Ala Ile  
85 90 95

Gly Leu Cys Ile Asp Leu Lys Thr Glu Ile Asn Phe Pro Ala Ser Met  
100 105 110

Arg Cys Glu Thr Val Thr Glu Gln Phe Phe Glu Leu Tyr His Leu Ala  
115 120 125

Ile Arg Leu Lys Glu Glu Pro Thr Ala Phe Arg Ala Asp Val Leu Leu  
130 135 140

Ser His Tyr Leu Gly Leu Gly Glu Leu Leu Cys Phe Ser Leu Glu Glu  
145 150 155 160



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Ile Gly His Glu Tyr Ile Glu Lys Asn Lys Ile Asn His Glu Arg Gln  
165 170 175

Ser Asn Gly Tyr  
180

<210> 78  
<211> 158  
<212> PRT  
<213> Bacillus anthracis  
  
<400> 78

Leu Ser Val Thr Lys Gly Val Cys Ile Asp Val Asp His Ser Asp Leu  
1 5 10 15

Leu His Glu Lys Val Glu Tyr Phe Leu Phe Pro Ala Lys Pro Ser His  
20 25 30

Tyr Tyr Val Ser Arg Phe Asn Arg Lys Gly Ala His Phe Gly Cys Tyr  
35 40 45

Gln Ala Glu Arg Phe Gln Ile Thr Glu Lys Glu Val Trp Thr Pro Glu  
50 55 60

Pro Gln Pro Asn Leu Pro Glu Leu Asn Thr Ser Leu Phe Tyr Arg Ala  
65 70 75 80

Gln Leu Ile Trp Arg Lys Lys Gly Tyr Lys Asp Lys Pro Leu Lys Asp  
85 90 95

Tyr Ile Val Gln Pro Arg Gly Lys His Cys Tyr Phe Trp His Asp Arg  
100 105 110

Glu Arg Lys Lys Phe Cys Gly Cys Phe Pro Leu His Trp Phe Thr Asp  
115 120 125

Phe Val Pro Val Gln Ser His His Ile Glu Glu Lys Thr Arg Glu Glu  
130 135 140

Val Lys Leu Leu Gln Arg Pro Asp Gly Gln Leu Ala Phe Phe  
145 150 155

<210> 79  
<211> 76  
<212> PRT  
<213> Bacillus anthracis  
  
<400> 79

Met Arg Val Ile Glu Ile Ser Trp Trp Ala Ile Ala Ile Gly Leu Tyr  
1 5 10 15

Leu Leu Ile Gly Val Ala Leu Leu Ile Trp Ile Ile Ala Thr Asp Ser  
Page 125

Trp Gly Ser Leu Phe Leu Tyr Pro Val Phe Ala Val Val Ile Val Leu  
35 40 45

Gly Trp Leu Pro Leu Met Ile Arg Ser Ile Val Gln Glu Ile Ser Lys  
50 55 60

Ala Ile His Lys Trp Lys Arg Lys Gln Lys Thr Glu  
65 70 75

<210> 80  
<211> 180  
<212> PRT  
<213> Bacillus anthracis

<400> 80

Met Asp Ile Lys Lys Leu Phe Ala Met Gln Asn Ile Leu Asp Lys Arg  
1 5 10 15

Val Leu Glu Ser Lys Asn Leu Ser Arg Gly Glu Val Phe Glu Phe Arg  
20 25 30

Ile Leu Ala Phe Leu Asp Glu Leu Gly Glu Cys Met Lys Glu Trp Arg  
35 40 45

Val Phe Lys Phe Trp Ser Asp Asp Arg Lys Pro Arg Thr Ser Ile Pro  
50 55 60

Thr Gly Glu Ile Ile Val Leu Asp Asp Gly Tyr Glu Val Glu Val Tyr  
65 70 75 80

Lys Asn Pro Leu Leu Glu Glu Tyr Val Asp Gly Leu His Phe Ala Ile  
85 90 95

Gly Leu Cys Ile Asp Leu Lys Thr Glu Ile Asn Phe Pro Ala Ser Met  
100 105 110

Arg Cys Glu Thr Val Thr Glu Gln Phe Phe Glu Leu Tyr His Leu Ala  
115 120 125

Ile Arg Leu Lys Glu Glu Pro Thr Ala Phe Arg Ala Asp Val Leu Leu  
130 135 140

Ser His Tyr Leu Gly Leu Gly Glu Leu Leu Cys Phe Ser Leu Glu Glu  
145 150 155 160

Ile Gly His Glu Tyr Ile Glu Lys Asn Lys Ile Asn His Glu Arg Gln  
165 170 175

Ser Asn Gly Tyr

<210> 81  
<211> 156  
<212> PRT  
<213> Bacillus anthracis

<400> 81

Met Ser Gly Cys Thr Ile Val Asn Val Lys Ile Asn Lys Gln Lys Arg  
1 5 10 15  
Gly Met Lys Asp Met Lys Trp Met Tyr Asn Leu Asp Ser Asn Asn Glu  
20 25 30  
Ile Trp Thr Ser Asp Lys Phe Glu Met Lys Glu Glu Ala Ile Gln Ala  
35 40 45  
Ala Leu Lys Asp Trp Thr Asp Lys Met Val Ala Asp Arg Ala Ala Val  
50 55 60  
Asp Asn Glu Phe Gln Ile Gly Gln Phe Lys Gln Tyr Ser Pro Trp Ile  
65 70 75 80  
Asn Ala Asp Val Leu Leu Asp Glu Leu Tyr Glu Arg Ala Thr Asp Glu  
85 90 95  
Cys Gly Glu Val Ala Glu Tyr Trp Leu Ser Gly Val Pro Met Asp Glu  
100 105 110  
Gly Glu Lys Leu Gln Glu Gln Ile Asn Lys Val Val Thr Glu Trp Leu  
115 120 125  
Lys Gly Ile Asn Glu His Pro Ser Phe Gly Ser Ile Glu Asn Ile Glu  
130 135 140  
Thr Ile Asp Ala Ser Lys Ile Glu Tyr Lys Glu Asn  
145 150 155

<210> 82  
<211> 310  
<212> PRT  
<213> Bacillus anthracis

<400> 82

Met Asp Cys Phe Lys Lys Gly Lys Phe Ile Pro Phe Pro Cys Ala Leu  
1 5 10 15  
Pro Ile Pro Glu Ala Gly Pro Thr Gly Pro Thr Gly Pro Pro Gly Ser  
20 25 30  
Ala Gly Gly Ser Thr Gly Pro Thr Gly Pro Thr Gly Pro Gln Gly Leu  
35 40 45

Gln Gly Ile Gln Gly Val Gln Gly Asn Pro Gly Thr Thr Gly Pro Gln  
50 55 60

Gly Ile Gln Gly Ile Gln Gly Ile Pro Gly Val Ser Gly Pro Ile Gly  
65 70 75 80

Pro Ile Gly Pro Thr Gly Ile Gln Gly Val Gln Gly Ile Gln Gly Phe  
85 90 95

Pro Gly Ile Pro Gly Pro Met Gly Pro Ile Gly Leu Thr Gly Pro Thr  
100 105 110

Gly Ile Gln Gly Ile Gln Gly Ile Gln Gly Val Gln Gly Ile Gln Gly  
115 120 125

Ile Gln Gly Asp Val Gly Pro Thr Gly Pro Gln Gly Ile Pro Gly Ile  
130 135 140

Pro Gly Leu Thr Gly Pro Thr Gly Ser Gln Gly Val Thr Gly Val Thr  
145 150 155 160

Gly Pro Ser Gly Gly Pro Pro Gly Pro Thr Gly Ala Thr Gly Pro Thr  
165 170 175

Gly Pro Ala Gly Gly Pro Pro Gly Pro Thr Gly Pro Thr Gly Pro Ala  
180 185 190

Gly Gly Pro Thr Gly Leu Thr Gly Pro Thr Gly Pro Thr Gly Pro Thr  
195 200 205

Gly Ile Gln Gly Ile Gln Gly Val Gln Gly Thr Gln Gly Ile Pro Gly  
210 215 220

Pro Thr Gly Pro Gln Gly Ile Gln Gly Val Gln Gly Leu Gln Gly Ile  
225 230 235 240

Pro Gly Ile Pro Gly Ser Met Gly Pro Thr Gly Leu Thr Gly Pro Thr  
245 250 255

Gly Leu Gln Gly Ile Gln Gly Ile Gln Gly Asn Pro Gly Pro Thr Gly  
260 265 270

Pro Phe Gly Pro Thr Gly Pro Thr Gly Leu Gln Gly Ile Gln Gly Leu  
275 280 285

Gln Gly Ile Gln Gly Ile Pro Gly Ser Asn Arg Thr Ser Arg Asn Pro  
290 295 300

Arg Ser Asn Arg Thr Cys

305

310

<210> 83  
<211> 129  
<212> PRT  
<213> Bacillus anthracis

<400> 83

Met Tyr Gln Thr Trp Lys Asn Leu Leu Asn Ser Ile Lys Lys Ile Leu  
1 5 10 15

Gln Ala Lys Leu Leu Val Lys Gly Arg Lys Leu Ala Tyr Phe Asp Leu  
20 25 30

Asn Gly Leu Trp Ile Ala Leu Asn Val Glu Glu Asp Ile Pro Arg Asn  
35 40 45

Glu Ile Lys Gln Ser Tyr Thr His Met Ala Phe Thr Val Thr Asn Glu  
50 55 60

Ala Leu Asp His Leu Lys Glu Val Leu Ile Gln Asn Asp Val Asn Ile  
65 70 75 80

Leu Pro Gly Arg Glu Arg Asp Glu Arg Asp Gln Arg Ser Leu Tyr Phe  
85 90 95

Thr Asp Pro Asp Gly His Lys Phe Glu Phe His Thr Gly Thr Leu Gln  
100 105 110

Asn Arg Leu Glu Tyr Tyr Lys Glu Asp Lys Lys His Met Thr Phe Tyr  
115 120 125

Ile

<210> 84  
<211> 163  
<212> PRT  
<213> Bacillus anthracis

<400> 84

Leu Leu Ala His Phe Pro Gln Lys Leu Phe Phe Phe Gly Gly Thr Asn  
1 5 10 15

Ser Gly Phe Gln Arg Ile Ala Gly Ser Pro Gly Ala Asp Ser Gln Asp  
20 25 30

Ile Pro Tyr Val Leu Gly Gly Ala Gly Ser Val Val Gly Leu Ser Ala  
35 40 45

Ser Ile Ser Ile Asn Asn Leu Pro Ile Gly Val Tyr Thr Ile Arg Val  
50 55 60

Cys Lys Asn Val Pro Ile Asn Leu Ala Ala Pro Gly Pro Gly Gln Val  
65 70 75 80

Ile Ser Thr Ile Ile Leu Thr Thr Thr Ala Val Ile Ser Gly Thr Ile  
85 90 95

Ile Leu Thr Ile Asn Pro Ser Asp Ile Gly Ala Gln Pro Val Arg Val  
100 105 110

Phe Asn Pro Asn Leu Val Ile Ala Pro Ala Thr Val Ala Trp Ser Ser  
115 120 125

Thr Ile Pro Gly Asp Ile Val Ala Arg Gly Asp Ala Met Ser Leu Phe  
130 135 140

Ile Thr Pro Gly Ile Thr Gln Asn Ala Val Tyr Thr Val Phe Leu His  
145 150 155 160

Thr Gly Asn

<210> 85  
<211> 56  
<212> PRT  
<213> Bacillus anthracis

<400> 85

Met Ile Val Lys Ala Thr Ile Lys Leu Glu Leu Asp Asp Ser Gln Lys  
1 5 10 15

Asn Trp Val Ser Tyr Val Arg Glu Gln Gly Gly Glu Glu Ala Val Phe  
20 25 30

His Tyr Leu Glu Glu Glu Val Gln Lys Lys Ile Glu Leu Ala Asp Phe  
35 40 45

Val Glu Met Lys Tyr Lys Asn Lys  
50 55

<210> 86  
<211> 191  
<212> PRT  
<213> Bacillus anthracis

<400> 86

Met Gln His Ile Pro Arg Tyr Tyr Tyr Gln Ser Gln Ser Pro Met Asp  
1 5 10 15

Ser Ile Trp Asn Asn Asn Asn Trp Ile Tyr Ala Trp Asn Pro Tyr Tyr  
20 25 30

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Tyr Asn Tyr Asn Asn Asn Ala Trp Asn Arg Asn Arg Asn Pro Tyr Cys  
35 40 45

Glu Asn Val Arg Leu Thr Asp Tyr Gly Ala Arg Pro Phe Val Leu Asn  
50 55 60

Ile Asn Gln Ala Thr Lys Gln Asn Asn Thr Tyr Arg Thr Ala Ile Trp  
65 70 75 80

Thr Gly Lys Asn Leu Gln Val Thr Leu Met Ser Ile Asn Val Gly Asp  
85 90 95

Asp Ile Gly Leu Glu Val His Pro Thr Thr Asp Gln Phe Ile Arg Ile  
100 105 110

Glu Glu Gly Gln Gly Leu Val Gln Met Gly Asp Asn Lys Asp Lys Leu  
115 120 125

Asp Phe Gln Glu Met Val Tyr Asp Asp Tyr Ala Ile Met Ile Pro Ala  
130 135 140

Gly Lys Trp His Asn Val Ile Asn Thr Gly Asn Thr Pro Leu Lys Ile  
145 150 155 160

Tyr Ala Ile Tyr Ala Pro Pro Glu His Pro Tyr Gly Thr Val His Glu  
165 170 175

Thr Lys Ala Ile Ala Met Ser Thr Glu Ala Asn Arg Tyr Tyr Tyr  
180 185 190

<210> 87  
<211> 101  
<212> PRT  
<213> Bacillus anthracis

<400> 87

Met Asp Met Ser Leu Val Gly Asn Leu Lys Glu Leu Gln Glu Lys Ala  
1 5 10 15

Ile Asp Glu Lys Val Leu Glu Phe Ala Glu Glu Met Glu Ile Val Ile  
20 25 30

Thr Lys Ser Ala Ala Ser Gly Tyr Ser Gly His Arg Tyr Lys Ile His  
35 40 45

Asn Glu Asn Pro Asn Arg His Met Met Cys Ser Lys Ile Phe Ile Glu  
50 55 60

Lys Leu Gln Glu Leu Leu Asp Gly Val Lys Val Glu Phe Lys Glu Glu  
65 70 75 80



Glu Lys Lys Asn Ile Leu Gly Gly Ser Tyr Tyr Glu His Tyr Ile Arg  
85 90 95

Phe Lys Trp Asn Asp  
100

<210> 88  
<211> 56  
<212> PRT  
<213> Bacillus anthracis

<400> 88

Met Ile Val Lys Ala Thr Ile Lys Leu Glu Leu Asp Asp Ser Gln Lys  
1 5 10 15

Asn Trp Val Ser Tyr Val Arg Glu Gln Gly Gly Glu Glu Ala Val Phe  
20 25 30

His Tyr Leu Glu Glu Glu Val Gln Lys Lys Ile Glu Leu Ala Asp Phe  
35 40 45

Val Glu Met Lys Tyr Lys Asn Lys  
50 55

<210> 89  
<211> 79  
<212> PRT  
<213> Bacillus anthracis

<400> 89

Met Thr Asn Phe Leu Leu Lys Ile Leu Phe Trp Arg Lys Gly Val Glu  
1 5 10 15

Arg Met Lys Thr Phe Asn Val Thr Phe Thr Glu Leu Lys Ile Tyr Glu  
20 25 30

Ala Val Ile Glu Ala Glu Ser Ala Glu Lys Ile Ile Asp Val Ile Lys  
35 40 45

His Leu Lys Arg Thr Glu Asp Asp Leu Val Asp Lys Gly Val Ile Ile  
50 55 60

Asn Glu Val Ser Glu Ile Asn Val Ser Lys Glu Gln Lys Phe Glu  
65 70 75

<210> 90  
<211> 101  
<212> PRT  
<213> Bacillus anthracis

<400> 90

Met Asp Met Ser Leu Val Gly Asn Leu Lys Glu Leu Gln Glu Lys Ala  
Page 132

Ile Asp Glu Lys Val Leu Glu Phe Ala Glu Glu Met Glu Ile Val Ile  
20 25 30

Thr Lys Ser Ala Ala Ser Gly Tyr Ser Gly His Arg Tyr Lys Ile His  
35 40 45

Asn Glu Asn Pro Asn Arg His Met Met Cys Ser Lys Ile Phe Ile Glu  
50 55 60

Lys Leu Gln Glu Leu Leu Asp Gly Val Lys Val Glu Phe Lys Glu Glu  
65 70 75 80

Glu Lys Lys Asn Ile Leu Gly Gly Ser Tyr Tyr Glu His Tyr Ile Arg  
85 90 95

Phe Lys Trp Asn Asp  
100

<210> 91  
<211> 135  
<212> PRT  
<213> Bacillus anthracis

<400> 91

Val Asn His His Leu Phe Asn Trp Leu Arg Asp Tyr Gln Lys Leu Glu  
1 5 10 15

Glu Asp Ile Ala Tyr Leu Glu Tyr Asn Leu Asp Lys Thr Lys Ala Glu  
20 25 30

Leu Arg Arg Trp Val Ser Gly Asp Leu Arg Glu Val Arg Leu Thr Ala  
35 40 45

Glu Ser Glu Gly Ala Lys Val Glu Asn Arg Ile Glu Ala Ile Glu Tyr  
50 55 60

Glu Leu Ala His Lys Met Asn Asp Met Tyr Lys Leu Lys Lys Leu Ile  
65 70 75 80

Ser Lys Phe Arg Gly Leu Glu Asn Gln Ile Leu Lys Leu Lys Tyr Val  
85 90 95

Asp Gly Met Thr Leu Glu Glu Ile Ala Glu Ala Val Asn Tyr Ser Ser  
100 105 110

Ser His Ile Lys Lys Lys His Ala Glu Leu Val Arg Leu Ile Lys Phe  
115 120 125

Val Glu Arg Glu Gly Val Ile

<210> 92  
 <211> 79  
 <212> PRT  
 <213> Bacillus anthracis

<400> 92

Met Thr Asn Phe Leu Leu Lys Ile Leu Phe Trp Arg Lys Gly Val Glu  
 1 5 10 15

Arg Met Lys Thr Phe Asn Val Thr Phe Thr Glu Leu Lys Ile Tyr Glu  
 20 25 30

Ala Val Ile Glu Ala Glu Ser Ala Glu Lys Ile Ile Asp Val Ile Lys  
 35 40 45

His Leu Lys Arg Thr Glu Asp Asp Leu Val Asp Lys Gly Val Ile Ile  
 50 55 60

Asn Glu Val Ser Glu Ile Asn Val Ser Lys Glu Gln Lys Phe Glu  
 65 70 75

<210> 93  
 <211> 74  
 <212> PRT  
 <213> Bacillus anthracis

<400> 93

Met Asp Val Gln Glu Leu Ser Arg Arg Leu Glu Asn Leu Glu His Lys  
 1 5 10 15

Val Leu Gln Val Glu Thr Lys Ala Asp Val Leu Asn Arg Thr Ala Ile  
 20 25 30

Gln Lys Gly Asp Lys Ile Lys Val Val Tyr Pro His Leu Gly Ile Gln  
 35 40 45

Gly Glu Tyr Leu Val Glu Lys Ile Asp Asn Gly Val Leu Glu Leu Val  
 50 55 60

Ala Glu Glu Thr Met Lys Lys Ile Gln Glu  
 65 70

<210> 94  
 <211> 135  
 <212> PRT  
 <213> Bacillus anthracis

<400> 94

Val Asn His His Leu Phe Asn Trp Leu Arg Asp Tyr Gln Lys Leu Glu  
 1 5 10 15

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Glu Asp Ile Ala Tyr Leu Glu Tyr Asn Leu Asp Lys Thr Lys Ala Glu  
20 25 30

Leu Arg Arg Trp Val Ser Gly Asp Leu Arg Glu Val Arg Leu Thr Ala  
35 40 45

Glu Ser Glu Gly Ala Lys Val Glu Asn Arg Ile Glu Ala Ile Glu Tyr  
50 55 60

Glu Leu Ala His Lys Met Asn Asp Met Tyr Lys Leu Lys Lys Leu Ile  
65 70 75 80

Ser Lys Phe Arg Gly Leu Glu Asn Gln Ile Leu Lys Leu Lys Tyr Val  
85 90 95

Asp Gly Met Thr Leu Glu Glu Ile Ala Glu Ala Val Asn Tyr Ser Ser  
100 105 110

Ser His Ile Lys Lys Lys His Ala Glu Leu Val Arg Leu Ile Lys Phe  
115 120 125

Val Glu Arg Glu Gly Val Ile  
130 135

<210> 95  
<211> 73  
<212> PRT  
<213> Bacillus anthracis

<400> 95

Leu Lys Lys Leu Ser Lys Gln Glu Leu Ala Ala Val Met Thr His Cys  
1 5 10 15

Ile Ser Thr Leu Gly Glu Gln Ile Val Asn Glu His Ile Asn Pro Gln  
20 25 30

Lys Leu Ala Gln Ala Ser Ala Leu His Asn Asp Leu Phe Asp Asn Thr  
35 40 45

Thr Pro Lys Glu Arg Arg Glu Ala Thr Ile Ser Leu Leu Gly Lys Ala  
50 55 60

Ile Asp Glu Phe Leu Glu Ser Lys Glu  
65 70

<210> 96  
<211> 74  
<212> PRT  
<213> Bacillus anthracis

<400> 96

PCT-US2005-009928\_Sequence Listing.txt as filed 10-3-05 in PCT  
Met Asp Val Gln Glu Leu Ser Arg Arg Leu Glu Asn Leu Glu His Lys  
1 5 10 15

Val Leu Gln Val Glu Thr Lys Ala Asp Val Leu Asn Arg Thr Ala Ile  
20 25 30

Gln Lys Gly Asp Lys Ile Lys Val Val Tyr Pro His Leu Gly Ile Gln  
35 40 45

Gly Glu Tyr Leu Val Glu Lys Ile Asp Asn Gly Val Leu Glu Leu Val  
50 55 60

Ala Glu Glu Thr Met Lys Lys Ile Gln Glu  
65 70

<210> 97  
<211> 134  
<212> PRT  
<213> Bacillus anthracis

<400> 97

Met Gly Lys Gly Tyr Phe Asn Lys Ala Val Cys Leu Val Cys Gly His  
1 5 10 15

Gln Asp Arg Val Asn His Pro Ser Lys Lys Glu Tyr Gln Glu Val Thr  
20 25 30

Val Cys Pro Glu Cys Asn Gly Ala Phe Val Asp Val Trp Lys Leu Gly  
35 40 45

Lys Tyr Lys Arg Asn Thr Gln Ser Asn Glu Glu Pro Leu Leu Thr Ile  
50 55 60

Thr Leu Thr Asp Ile Asp Ala Lys Pro Ile Val His Tyr Lys Gly Glu  
65 70 75 80

Gln Ile Asp Arg Lys Leu Arg Val Thr Phe Asp Trp Glu Ser Gln Ser  
85 90 95

Ile Asp Lys Ile Asn Arg Thr Tyr Ile His Ile Glu His Val Pro Ala  
100 105 110

Asp Asn Lys Arg Leu Asn Thr Glu Thr Ile Gln His Asn His Pro Ile  
115 120 125

Ala Asn Lys Glu Gln Val  
130

<210> 98  
<211> 73  
<212> PRT  
<213> Bacillus anthracis

<400> 98

Leu Lys Lys Leu Ser Lys Gln Glu Leu Ala Ala Val Met Thr His Cys  
1 5 10 15

Ile Ser Thr Leu Gly Glu Gln Ile Val Asn Glu His Ile Asn Pro Gln  
20 25 30

Lys Leu Ala Gln Ala Ser Ala Leu His Asn Asp Leu Phe Asp Asn Thr  
35 40 45

Thr Pro Lys Glu Arg Arg Glu Ala Thr Ile Ser Leu Leu Gly Lys Ala  
50 55 60

Ile Asp Glu Phe Leu Glu Ser Lys Glu  
65 70

<210> 99

<211> 63

<212> PRT

<213> Bacillus anthracis

<400> 99

Met Asn Gly Phe Asn Lys Ile Val Asn Asp Met Gln Asn Glu Gln Val  
1 5 10 15

Gly Asn Ala Met Leu Asp Phe Ala Leu Ala Ala Lys Met Met Phe Ala  
20 25 30

Ala Phe Thr Gln Phe Lys Glu Ala Gly Phe Asn Glu Glu Gln Ser Phe  
35 40 45

Glu Leu Thr Arg Glu Ile Leu Ile Asp Ser Leu Ser Lys Asn Gln  
50 55 60

<210> 100

<211> 134

<212> PRT

<213> Bacillus anthracis

<400> 100

Met Gly Lys Gly Tyr Phe Asn Lys Ala Val Cys Leu Val Cys Gly His  
1 5 10 15

Gln Asp Arg Val Asn His Pro Ser Lys Lys Glu Tyr Gln Glu Val Thr  
20 25 30

Val Cys Pro Glu Cys Asn Gly Ala Phe Val Asp Val Trp Lys Leu Gly  
35 40 45

Lys Tyr Lys Arg Asn Thr Gln Ser Asn Glu Glu Pro Leu Leu Thr Ile  
50 55 60

Thr Leu Thr Asp Ile Asp Ala Lys Pro Ile Val His Tyr Lys Gly Glu  
65 70 75 80

Gln Ile Asp Arg Lys Leu Arg Val Thr Phe Asp Trp Glu Ser Gln Ser  
85 90 95

Ile Asp Lys Ile Asn Arg Thr Tyr Ile His Ile Glu His Val Pro Ala  
100 105 110

Asp Asn Lys Arg Leu Asn Thr Glu Thr Ile Gln His Asn His Pro Ile  
115 120 125

Ala Asn Lys Glu Gln Val  
130

<210> 101  
<211> 84  
<212> PRT  
<213> Bacillus anthracis

<400> 101

Met Gln Val Tyr Cys Ser Glu Cys Asp Lys Ser Tyr Asp Met Gln Pro  
1 5 10 15

Gln Val Thr Gln Leu Pro Asn Arg Ile Glu Lys Cys Phe Phe Ile Cys  
20 25 30

Pro His Cys Asn His Glu His Ile Ala Ala Tyr Val Asn Asp Lys Ile  
35 40 45

Arg Lys Tyr Gln Ala Asp Ile Ala Lys Cys His Glu Arg Ile Asn Lys  
50 55 60

Lys Asn Leu Ala Ile Glu Asp Glu Met Lys Arg Leu Arg Lys Arg Phe  
65 70 75 80

Asp Arg Arg Lys

<210> 102  
<211> 63  
<212> PRT  
<213> Bacillus anthracis

<400> 102

Met Asn Gly Phe Asn Lys Ile Val Asn Asp Met Gln Asn Glu Gln Val  
1 5 10 15

Gly Asn Ala Met Leu Asp Phe Ala Leu Ala Ala Lys Met Met Phe Ala  
20 25 30



Ala Phe Thr Gln Phe Lys Glu Ala Gly Phe Asn Glu Glu Gln Ser Phe  
35 40 45

Glu Leu Thr Arg Glu Ile Leu Ile Asp Ser Leu Ser Lys Asn Gln  
50 55 60

<210> 103  
<211> 63  
<212> PRT  
<213> Bacillus anthracis

<400> 103

Met Glu Gly Gln Glu Leu Thr Leu Glu Lys Lys Asp Ser Ile Tyr Leu  
1 5 10 15

Arg Pro Arg Tyr Pro His Lys Ile Asp Ala Ser Lys Ile Lys Ser Leu  
20 25 30

Lys Asp Val Ile Lys Ile Leu Gly Leu Met Asp Ile Arg Leu Asp Asp  
35 40 45

Lys Ala Val Ile Gly Leu Glu His Leu Ile Glu Lys Glu Glu Glu  
50 55 60

<210> 104  
<211> 84  
<212> PRT  
<213> Bacillus anthracis

<400> 104

Met Gln Val Tyr Cys Ser Glu Cys Asp Lys Ser Tyr Asp Met Gln Pro  
1 5 10 15

Gln Val Thr Gln Leu Pro Asn Arg Ile Glu Lys Cys Phe Phe Ile Cys  
20 25 30

Pro His Cys Asn His Glu His Ile Ala Ala Tyr Val Asn Asp Lys Ile  
35 40 45

Arg Lys Tyr Gln Ala Asp Ile Ala Lys Cys His Glu Arg Ile Asn Lys  
50 55 60

Lys Asn Leu Ala Ile Glu Asp Glu Met Lys Arg Leu Arg Lys Arg Phe  
65 70 75 80

Asp Arg Arg Lys

<210> 105  
<211> 95  
<212> PRT  
<213> Bacillus anthracis

<400> 105

Leu Lys Arg Arg Lys Asn Lys Met Ala Asn Asn Lys Leu Ile Ile Glu  
1 5 10 15

Val Thr Ala Asp Thr Thr Glu Ala Leu Glu Gly Ile Lys Glu Val Thr  
20 25 30

Glu Ala Ala Asn Glu Cys Ala Asp Ala Leu Asp Lys Leu Glu Lys Ile  
35 40 45

Met Asp Lys Phe Thr Asn Arg Ser Asp Thr Val Glu Leu Tyr Cys Glu  
50 55 60

Gly Lys Leu Leu Ser Lys Ser Thr Val Asn His Thr Ala Asp Ser Ile  
65 70 75 80

Gln Cys Arg Ile Ile Lys Gly Glu Glu Leu Gly Gly Ser Glu Arg  
85 90 95

<210> 106

<211> 63

<212> PRT

<213> Bacillus anthracis

<400> 106

Met Glu Gly Gln Glu Leu Thr Leu Glu Lys Lys Asp Ser Ile Tyr Leu  
1 5 10 15

Arg Pro Arg Tyr Pro His Lys Ile Asp Ala Ser Lys Ile Lys Ser Leu  
20 25 30

Lys Asp Val Ile Lys Ile Leu Gly Leu Met Asp Ile Arg Leu Asp Asp  
35 40 45

Lys Ala Val Ile Gly Leu Glu His Leu Ile Glu Lys Glu Glu Glu  
50 55 60

<210> 107

<211> 127

<212> PRT

<213> Bacillus anthracis

<400> 107

Met Lys Lys Pro Leu Arg Pro Cys Cys Glu Phe His Cys Tyr Asn Leu  
1 5 10 15

Thr Arg Glu Arg Tyr Cys Glu Glu His Arg Tyr Lys Glu Lys Glu Thr  
20 25 30

Gln Gln Asp Lys Asn Arg Tyr Tyr Asp Arg Phe Lys Arg Asp Lys Glu  
35 40 45

Ser Thr Ala Phe Tyr Arg Ser Lys Ala Trp Glu Arg Leu Arg Glu Gln  
50 55 60

Ala Leu Met Arg Asp Lys Gly Leu Cys Leu His Cys Lys Asn Asn Arg  
65 70 75 80

Lys Ile Lys Val Ala Asp Met Val Asp His Ile Ile Pro Ile Lys Val  
85 90 95

Asp Pro Ser Leu Lys Leu Lys Leu Glu Asn Leu Gln Ser Leu Cys Asn  
100 105 110

Pro Cys His Asn Arg Lys Thr Ala Glu Asp Lys Lys Lys Tyr Gly  
115 120 125

<210> 108

<211> 95

<212> PRT

<213> Bacillus anthracis

<400> 108

Leu Lys Arg Arg Lys Asn Lys Met Ala Asn Asn Lys Leu Ile Ile Glu  
1 5 10 15

Val Thr Ala Asp Thr Thr Glu Ala Leu Glu Gly Ile Lys Glu Val Thr  
20 25 30

Glu Ala Ala Asn Glu Cys Ala Asp Ala Leu Asp Lys Leu Glu Lys Ile  
35 40 45

Met Asp Lys Phe Thr Asn Arg Ser Asp Thr Val Glu Leu Tyr Cys Glu  
50 55 60

Gly Lys Leu Leu Ser Lys Ser Thr Val Asn His Thr Ala Asp Ser Ile  
65 70 75 80

Gln Cys Arg Ile Ile Lys Gly Glu Glu Leu Gly Gly Ser Glu Arg  
85 90 95

<210> 109

<211> 127

<212> PRT

<213> Bacillus anthracis

<400> 109

Met Lys Lys Pro Leu Arg Pro Cys Cys Glu Phe His Cys Tyr Asn Leu  
1 5 10 15

Thr Arg Glu Arg Tyr Cys Glu Glu His Arg Tyr Lys Glu Lys Glu Thr  
20 25 30

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Gln	Gln	Asp	Lys	Asn	Arg	Tyr	Tyr	Asp	Arg	Phe	Lys	Arg	Asp	Lys	Glu
		35					40					45			
Ser	Thr	Ala	Phe	Tyr	Arg	Ser	Lys	Ala	Trp	Glu	Arg	Leu	Arg	Glu	Gln
	50					55				60					
Ala	Leu	Met	Arg	Asp	Lys	Gly	Leu	Cys	Leu	His	Cys	Lys	Asn	Asn	Arg
65					70					75					80
Lys	Ile	Lys	Val	Ala	Asp	Met	Val	Asp	His	Ile	Ile	Pro	Ile	Lys	Val
				85					90					95	
Asp	Pro	Ser	Leu	Lys	Leu	Lys	Leu	Glu	Asn	Leu	Gln	Ser	Leu	Cys	Asn
			100					105					110		
Pro	Cys	His	Asn	Arg	Lys	Thr	Ala	Glu	Asp	Lys	Lys	Lys	Tyr	Gly	
		115					120					125			